
COMPOSITE ANALYTICAL LABORATORY REPORT

ON THE

Quantitative Analysis of Fluorochemicals in Environmental Samples

**REPORT NO. FACT GEN-021, GEN-024, GEN-030, GEN-033
LRN—W2491, W2845, W3197, E00-1386**

ANALYTICAL STUDY INITIATION

GEN021: 08/25/99
GEN024: 10/12/99
GEN030: 12/13/99
GEN033: 03/14/00

**Exhibit
2814**

State of Minnesota v. 3M Co.,
Court File No. 27-CV-10-28862

3M_MN01665997

2814.0001

TABLE OF CONTENTS

Analytical Study Personnel and Contributors	1
Introduction	2
Purpose	2
Test and Control Article	2
Sample Collection and Analysis	3
Sample Receipt and Maintenance	3
Chemical Characterization	3
Procurement	3
Method Summaries	3
Preparatory and Analytical Methods	4
Analytical Equipment	5
Data Summary, Analyses, and Results	6
Summary of Quality Control Analyses Results	6
Summary of Sample Results	7
Data Quality Objectives	8
Statement of Conclusion	9
References	9
Attachments	10
Report Signature	10

LIST OF TABLES

Table 1. Description of Samples, by Study	2
Table 2. Procurement Information for Reference Materials in the Analysis of Environmental Samples	3
Table 3. Ions Monitored in the Analyses of Extracts of Groundwater	6
Table 4. Range of LOQs for Sera, by Study	8
Table 5. Range of LOQs for Liver and Other Tissues, by Study	8

ANALYTICAL STUDY PERSONNEL AND CONTRIBUTORS

Analytical Chemistry Laboratories

3M Environmental Technology and Safety Services (ET&SS)
3M Environmental Laboratory
Fluorine Analytical Chemistry Team (FACT)
2-3E-09
935 Bush Avenue
St. Paul, MN 55106

Kris Hansen, Ph.D., *PAI*
Harold Johnson, *Analytical Chemist*
Lisa A. Clemen, *Analytical Chemist*
Mark Ellefson, *Analytical Chemist*

Sponsor

3M ET&SS
Building 2-3E-09
St. Paul, MN 55133

Dale Bacon, *Sponsor Representative*

INTRODUCTION

Purpose

The purpose of this composite report is to provide a summary of the analytical data collected for 3M studies Gen-021, Gen-024, Gen-030, and Gen-033. All of the samples included in these studies are tissue samples collected from fish, birds, mammals, and amphibians; Dr. John Geisy of Michigan State University has supplied all samples to 3M. These analyses have been conducted to support studies designed by Dr. Geisy.

The target analytes for these four studies were perfluorooctane sulfonate (PFOS; CAS# 2795-39-3), perfluorooctanesulfonylamide (PFOSA; CAS# 754-91-6), perfluorooctanoate (PFOA or POAA; CAS# 3825-26-1), and perfluorohexane sulfonate (PFHS; no CAS# available).

Due to the variety of matrices analyzed (with respect to both species and tissues), and due to evolving analytical methods, some analytical data quality objectives, such as the limit of quantitation (LOQ) were quite variable. A summary of the achieved LOQ (by specie, tissue and study number) is presented in Table 2 of this report. The stated data quality is based on results of data collection quality controls, sample prep quality controls, and recovery of target analytes from prepared matrix spike samples. More specific data quality objectives and parameters for these analytical studies are outlined later in this report.

Test and Control Article

The test articles for each study consisted of various tissues from various species and are listed below, in Table 1. For all studies, the control article consisted of rabbit sera and rabbit liver, as appropriate. Rabbit tissues were chosen as the control articles because previous studies have indicated very low levels of endogenous fluorochemicals in these matrices. Samples of the control articles were provided by the 3M Environmental Laboratory.

This report does not include details for the collection of the test articles; these details should be obtained from Dr. Geisy.

Table 1. Description of Samples, by Study

STUDY NUMBER	SERA/PLASMA/BLOOD	LIVER	OTHER
Gen-021	Cormorant Blood, Caspian Seal Blood, Sea Otter Blood	California Sea Lion, Elephant Seal, Harbor Seal, Gozzi, Mink, River Otter, Sea Otter, Turtle	Sea Otter Brain, Sea Otter Kidney
Gen-024	Albatross sera, Albatross plasma, Cormorant plasma, Herring Gull Plasma, Bald Eagle plasma, Cormorant blood, Herring Gull blood	Loon, Brown Pelican, Albatross	Albatross kidney, Cormorant yolk, Gull yolk
Gen-030	Northern Fur Seal blood (juvenile, sub-adult, adult), Polar Bear blood, Stellar Sea Lion blood	Northern Fur Seal, Polar Bear, Mink, Map Turtle, Terrapin, Tuna, Green Frog, Chinook Salmon, Lake Whitefish, Brown Trout	Carp body, Frog muscle, Frog body, Green Frog eggs, Lake Whitefish eggs, Brown Trout eggs, Carp muscle, Chinook Salmon muscle, Lake Whitefish muscle, Brown Trout muscle
Gen-033	None Submitted	Mink, Baikal Seal, Ganges Dolphin, Cormorant (adult and juvenile), Bottlenose Dolphin, Striped Dolphin, Weddell Seal, Swordfish, Tuna, Blacktailed Gull	None Submitted

Following analysis, extracts generated from these samples have been retained in cold storage.

Sample Collection and Analysis

Tissue samples were submitted to the Environmental Laboratory- Fluorine Analytical Chemistry Team by Kurunthachalan Kannan of Michigan State University. Details of the sample receipt are documented on the chain of custody forms located in appendices of this report.

SAMPLE RECEIPT AND MAINTENANCE

Samples were received in the Environmental Lab cold or frozen on the following dates: Gen-021 (8/24/99), Gen-024 (10/11/99), Gen-030 (12/13/99), and Gen-033 (3/13/00). Sample receipt, identification, and chain of custody information are located in the study folder for each report; the folders are located in the 3M archives.

The sample extracts will be maintained in cold storage at the 3M Environmental Laboratory until the quality of preparation no longer affords preservation.

CHEMICAL CHARACTERIZATION

The target analytes characterized in the samples include PFOS, PFOSA, PFOA, and PFHS. Procurement details of the reference standards used for analysis are summarized below.

Procurement

Table 2. Procurement Information for Reference Materials in the Analysis of Environmental Samples

REFERENCE MATERIAL	LOT NUMBER	SOURCE
PFOS (potassium salt)	171	3M ICP/PCP Division
PFOSA	Gen-021: L-2353; all others: L-15709	3M Specialty Chemicals (R. Buckanin)
PFHS (potassium salt)	NB116638-16	3M Specialty Chemicals (G. Moore)
PFOA (ammonium salt)	Gen-024: 245; all others: commercial	Gen-024: 3M Specialty Chemicals; all others: Aldrich

Full chemical characterization studies, including purity and stability determination, have not been completed at this time. Upon completion of these studies, a report will be archived in the 3M Environmental Lab.

METHOD SUMMARIES

Following is a brief description of the methods used during this analytical study by the 3M Environmental Laboratory. Copies of the actual methods used for these studies are located in attachment H.

PREPARATORY AND ANALYTICAL METHODS

- ETS-8-004.1, "Extraction of PFOS or Other Fluorochemical Compounds from Serum for Analysis using HPLC-Electrospray Mass Spectrometry" with some modifications, described below.

Because the matrices were so variable and sample size extremely limited, it was not possible to prepare extracted standard curves. All extracts were evaluated versus unextracted standard curves. When sample size permitted, two matrix spikes were prepared in each tissue sample from each specie tested to provide some level of extraction efficiency determination.

For some samples, less than 1 mL of sample was available. For these samples, the available volume was extracted according to the method with the exception that the final volume of extraction solvent was adjusted to match the volume of the initial sample.

This method was used for the extraction of sera, plasma, and whole blood samples.

- ETS-8-005.1, "Analysis of PFOS or Other Fluorochemical Compounds in Serum Extracts Using HPLC-Electrospray Mass Spectrometry" with some modifications, described below.

Because the matrices were so variable and sample size extremely limited, it was not possible to prepare extracted standard curves. All extracts were evaluated versus unextracted standard curves; as a result, all sample concentrations were adjusted by a factor of 1.25 to adjust for the removal of 4/5 of the MTBE from the extract. The factor is unnecessary when an extracted curve is used for evaluation.

- ETS-8-006, "Analysis of PFOS or Other Fluorochemical Compounds in Liver Extracts using HPLC-Electrospray Mass Spectrometry" with some modifications, described below.

Because the matrices were so variable and sample size extremely limited, it was not possible to prepare extracted standard curves. All extracts were evaluated versus unextracted standard curves. When sample size permitted, two matrix spikes were prepared in each tissue sample from each specie tested to provide some level of extraction efficiency determination.

For some samples, less than 1 g of sample (as called for in the method) was available. For these samples, the available mass of tissue was extracted according to the method.

Samples of kidney, brain, egg, and muscle were extracted by this method.

- ETS-8-007, "Extraction of PFOS or Other Fluorochemical Compounds from Liver for Analysis using HPLC-Electrospray Mass Spectrometry" with some modifications, described below.

Because the matrices were so variable and sample size extremely limited, it was not possible to prepare extracted standard curves. All extracts were evaluated versus unextracted standard curves; as a result, all sample concentrations were adjusted by a factor of 1.25 to adjust for the removal of 4/5 of the MTBE from the extract. The factor is unnecessary when an extracted curve is used for evaluation.

For Gen-030 and Gen-033 only: Due to the lack of excess test material for method development, all samples determined to contain greater than 0.015 µg/g of PFOS were subject to an additional PFOS verification process. Each sample was analyzed separately with respect to the 499 → 99 transition and the 499 → 80 transitions. The quantitative results

obtained from each transition analysis were compared. When these results agreed to within 30%, the identity of PFOS was confirmed (see Reference 1). Those samples where the identity of PFOS could not be confirmed are noted in the data table.

In Gen-021, no PFHS standard was available. In these samples, qualitative determination of PFHS was conducted based on reasonable retention time and a known PFHS transition (399 → 99).

Specific instrumental parameters are available in appendix I-L of this report, stored in the 3M Environmental Lab archives.

ANALYTICAL EQUIPMENT

For HPLC-Electrospray Tandem Mass Spectrometry:

Liquid Chromatograph: Hewlett-Packard® Series 1100 Liquid Chromatograph system

Analytical column:

1×30 mm C18 Betasil™

Column temperature: 30 degrees C

Cycle Time: 10 minutes

Mobile phase components:

Component A: 2mM ammonium acetate

Component B: Methyl alcohol

Flow rate: 300 µL/min

Injection volume: 10 µL

Solvent Gradient:

Time (min)	%B
0	10
1	10
5.5	95
7.5	95
8	10
10	10

For Detection:

Mass Spectrometer: Micromass® API/Mass Spectrometer Quattro Ultima Triple Quadrupole system or Micromass® API/Mass Spectrometer Quattro II Triple Quadrupole system

Acquisition Mode: MRM (refer to Table 3)

Software: Mass Lynx™ 3.3

Mode: Electrospray Negative

Source Block Temperature: 125-150°C

Source: Z-spray

Table 3. Ions Monitored in the Analyses of Extracts of Groundwater

TARGET ANALYTE	PRIMARY ION (amu)	PRODUCT ION (amu)
PFOS	499.0	80, 99*
PFOSA	498	78
PFOA	413	169
PFHS	399	99

* Indicates the ion used for quantitation

Refer to the analytical methods and equipment logs found in the raw data for details on the actual analytical equipment settings used in the present study. These settings may have varied somewhat during actual data collection. However, slight variations in the instrument settings will not adversely affect the quality of the data. Exact settings during all phases of data collection are recorded and presented in the appendix of this report.

DATA SUMMARY, ANALYSES, AND RESULTS

Summary of Quality Control Analyses Results

- **Standard Curves:** The coefficient of determination (r^2) for all 1/X weighted curves bracketing useable data was ≥ 0.982 . High or low curve points may have been excluded to provide a better fit over the linear range appropriate to the data. High or low curve points were deactivated if the calculated concentration varied from the theoretical concentration by more than 30%. Acceptable data was evaluated versus a standard curve containing at least 5 points. All actions are acceptable and are documented in specific data sets. All standard curves used to evaluate quantitative data were acceptable.
- **Continuing Calibration Verifications:** On average, one calibration check is analyzed for every five samples. Acceptable data is bracketed by calibration checks quantitated to be within 30% of the theoretical value, evaluated at least every ten samples. All quantitative data is bracketed by acceptable calibration checks, as required.
- **Blanks:** Extraction blanks were compliant if no target analyte was detected above the limit of quantitation (LOQ) for a specific analyte. In this study, extraction blanks were often higher than low curve points. Because analyte levels in the blank are used to determine the LOQ, by default, all blanks were determined to be below the limit of quantitation for the compounds of interest.
- **Internal Standards:** Internal standard response was monitored in Gen-030 and Gen-033 only. Internal standard response was required to be within $\pm 50\%$ of the theoretical value. If samples showed an internal standard response that deviated more than $\pm 50\%$, the samples were reanalyzed. If the deviant IS response was confirmed, the analyte data was reported, but noted in the data table.

Summary of Sample Results

- **GEN-021:**
 - PFOS was detected in at least one sample from the following matrices: California Sea Lion liver, Harbor Seal liver, Gozzi liver, Mink liver, River Otter liver, Turtle liver, Cormorant blood, Otter blood, and Caspian Seal blood.
 - PFOSA was tentatively identified in at least one sample from the following matrices: California Sea Lion liver, River Otter liver, Sea Otter liver, Sea Otter brain, and Otter blood.
 - PFOA was tentatively identified in at least one sample from the following matrices: California Sea Lion liver and Caspian Seal blood.
 - PFHS was tentatively identified in at least one sample from the following matrices: California Sea Lion liver, Gozzi liver, Mink liver, River Otter liver, Sea Otter liver, Turtle liver, Cormorant blood, Caspian Seal blood, and Otter blood.
- **GEN-024:**
 - PFOS was detected identified in at least one sample from the following matrices: Albatross plasma, Albatross sera, Cormorant plasma, Cormorant blood, Herring Gull plasma, Herring Gull blood, Bald Eagle plasma, Loon liver, Albatross liver, Brown Pelican liver, Albatross kidney, Cormorant yolk, and Gull yolk.
 - PFOSA was tentatively identified in at least one sample from the following matrices: Cormorant blood, Bald Eagle plasma, Loon liver, Brown Pelican liver, and Albatross liver.
 - PFOA was tentatively identified in at least one sample from the following matrices: Cormorant blood, Albatross liver, Cormorant yolk, and Gull yolk.
 - PFHS was tentatively identified in at least one sample from the following matrices: Herring Gull plasma and Bald Eagle plasma, Loon liver, Albatross liver, Brown Pelican liver, Albatross kidney, Cormorant yolk, and Gull yolk.
- **GEN-030:**
 - PFOS was detected in at least one sample from the following matrices: Polar Bear blood, Polar Bear liver, Mink liver, Northern Fur Seal liver, Map Turtle liver, Tuna liver, Green Frog liver, Chinook Salmon liver, Lake Whitefish liver, Brown Trout liver, Whole Carp, Frog muscle, Lake Whitefish eggs, Brown Trout eggs, Carp muscle, Chinook Salmon muscle, Lake Whitefish muscle, and Brown Trout muscle.
 - PFOSA was tentatively identified in at least one sample from the following matrices: Mink liver.
 - PFOA was not tentatively identified in any sample analyzed.
 - PFHS was not tentatively identified in any sample analyzed.
- **GEN-033:**
 - PFOS was detected in at least one sample from the following matrices: Mink liver, Baikal Seal liver, Cormorant liver, Bottle Nosed Dolphin liver, Ganges Dolphin liver, Striped Dolphin liver, Swordfish Liver, Tuna liver, and Black Tailed Gull liver.
 - PFOSA was tentatively identified in at least one sample from the following matrices: Mink liver, Cormorant liver, and Bottle Nosed Dolphin liver.

- PFOA was tentatively identified in at least one sample from the following matrices: Cormorant liver.
- PFHS was tentatively identified in at least one sample from the following matrices: Mink liver, Striped Dolphin liver, and Swordfish Liver.

Appendices contain data summary tables.

DATA QUALITY OBJECTIVES

No circumstances existed during the present study that would have affected the quality or integrity of the data. The data quality objectives (DQOs) followed during the present are indicated below.

- **Linearity:** The coefficient of determination (r^2) of the standard curve was equal to or greater than 0.985 with at least 5 active points using a linear regression curve with 1/x weighting.
- **Instrument Quantitation Limit (IQL):** The IQL is equal to the lowest acceptable standard in the calibration curve (acceptable standard is defined as a standard within 30% of the theoretical value). As this value is not useful in consideration of the sample data, the IQL was not specifically determined or stated for every study.
- **Limits of Quantitation (LOQ):** The LOQ is equal to the lowest acceptable standard in the calibration curve (defined as a standard within 30% of the theoretical value), and is at least two times the analyte peak area detected in the extraction blanks. The LOQ may vary due to the amount of sample available for analysis (particularly for samples extracted according to ETS-8-006) or to day-to-day variations in the analytical system. The ranges of LOQs for various tissues are listed in Table 4 (sera, plasma, and blood) and Table 5 (liver, kidney, muscle, egg, and brain).

Table 4. Range of LOQs for Sera, by Study

ANALYTE	GEN-021	GEN024	GEN-030	GEN-033
PFOS	0.0116 µg/mL	0.00116 µg/mL	0.0029-0.0579 µg/mL	NA
PFOSA	0.00625µg/mL	0.00626 µg/mL	0.000625 µg/mL	NA
PFHS	NA	0.00114 µg/mL	0.00114 µg/mL	NA
PFOA	0.00599 µg/mL	0.0299 µg/mL	0.00240-0.00958 µg/mL	NA

Table 5. Range of LOQs for Liver and Other Tissues, by Study

ANALYTE	GEN-021	GEN024	GEN-030	GEN-033
PFOS	0.0348 µg/g	0.0348 µg/g	0.00696-0.0696 µg/g	0.00696- 0.0694µg/g
PFOSA	0.0375 µg/g	0.00750 µg/g	0.0188 µg/g	0.0376 µg/g
PFHS	NA	0.00683 µg/g	0.00683-0.0342 µg/g	0.00683 µg/g
PFOA	0.0359 µg/g	0.180 µg/g	0.0180-0.0719 µg/g	0.00719-0.0718 µg/g

NA = not applicable

- **Duplicate/acceptable precision (extraction):** Spikes conducted on samples of control tissues were reproducible to within 15%
- **Quality Control Response:** A continuing calibration verification (CCV) was analyzed every 5–10 samples. Acceptable CCV response was within $\pm 30\%$ of the theoretical value. No more than 10 samples were analyzed between acceptable CCVs.
- **Spike/acceptable recoveries:** Due to the number of different matrices analyzed, there was great variability in spike recoveries. For any given matrix (specie and tissue), spike recoveries within 70–130% of the expected concentration indicate quantitative data (good to $\pm 30\%$); spike recoveries between 50–150% indicate semi-quantitative data for that matrix (good to $\pm 50\%$). Spike recoveries outside of this range indicate that sample data should be used for qualitative purposes only. Due to sample limitations, matrix spike studies were not conducted for all matrices. For PFOS analyses, sample data that is not supported by matrix spike studies should be considered for qualitative purposes only. Since no identity verification experiments were performed for PFOA, PFHS, and PFOSA, for these analytes, all analyses that are not supported by matrix spike studies should be considered to provide unconfirmed qualitative data only.
- **Use of Internal Standards:** Tetrahydro-perfluorooctane sulfonate (THPFOS) was spiked into the extracts post-extraction and used as an internal standard for samples in Gen-030 and Gen-033. For all samples in these studies, THPFOS levels were monitored to verify the analytical soundness of the data. THPFOS levels that were determined to be deviant from expected values by more than $\pm 50\%$ were reanalyzed. If the deviant THPFOS levels were confirmed, analyte levels were reported but are noted in the results table.
- **Use of confirmatory methods:** Given the selectivity of the analytical tool used (HPLC-ESMSMS) and lack of a viable alternative for analysis, no confirmatory methods were used.
- **Demonstration of specificity:** Specificity was demonstrated by chromatographic retention time (matched to standards to within 3%) and the response of at least one characteristic product ion arising from collisions of an analyte-specific parent ion.

Assuming spike recovery studies form a suitable indication of endogenous analyte recovery, matrix spike studies have been used as an indicator of data quality (see above). The validity of this assumption has not been verified by other techniques.

STATEMENT OF CONCLUSION

Under the conditions of the present studies, the presence of fluorochemicals was observed in the quantitative analysis of a selection of environmental matrices.

REFERENCES

- 1) "Acceptance Criteria for Ultratrace HPLC-Tandem Mass Spectrometry: Quantitative and Qualitative Determination of Sulfonylurea Herbicides in Soil"; Li, L.Y.; Campbell, D.A.; Bennet, P.K.; Henion, J.; *Anal. Chem.*, **68** (19), 3397-3404, 1996

ATTACHMENTS

- Attachment A: Gen-021 Sera/Plasma/Blood Results
 - Attachment B: Gen-021 Liver/Miscellaneous Results
 - Attachment C: Gen-024 Sera/Plasma/Blood Results
 - Attachment D: Gen-024 Liver/Miscellaneous Results
 - Attachment E: Gen-030 Sera/Plasma/Blood Results
 - Attachment F: Gen-030 Liver/Miscellaneous Results
 - Attachment G: Gen-033 Liver Results
 - Attachment H: Analytical Methods
 - Attachment I – L (additional bound document available in the 3M Environmental Lab archives): Analytical Details for Gen-021, Gen-024, Gen-030, and Gen-033
-

REPORT SIGNATURE

 5/10/00
Kristen J. Hansen, Ph.D., Principal Analytical Investigator Date

 5/10/2000
Dale L. Bacon, Sponsor Representative Date

A

Study:
 Product Number(Test Substance):
 Matrix:
 Method/Revision:
 Analytical Equipment System Number:
 Instrument Software/Version:
 Filename:
 R-Squared Value:
 Slope:
 Y-Intercept:
 Dates of Extraction/Analyst:
 Dates of Analysis/Analyst:
 Date of Data Reduction/Analyst:

GEN021 Various Matrices from MSU
 None
 Blood
 ETS-8-4.1 & ETS-8-5.1 using unextracted curves
 Amelia 062498
 Masslynx 3.2
 See list to right
 See Attachments
 See Attachments
 See Attachments
 08/25/99 MCH/KK/SAL/SEE
 08/26/99, 08/27/99, 08/28/99, 12/09/99 MEE/IAS
 08/27/99, 08/30/99, 08/31/99, 12/10/99, 01/20/00 MEE/IAS

Sample Data

BLOOD

Group Dose	Sample #	Concentration of PFOS ug/mL	Mean PFOS ug/mL	RSD Std. Dev.	Concentration of PFOSA ug/mL	Mean PFOSA ug/mL	RSD Std. Dev.
Method Blk	H2O Blk-1 8/25/99	<LOQ (0.0116)			<LOQ (0.00625)		
	H2O Blk-2 8/25/99	<LOQ (0.0116)			<LOQ (0.00625)		
	H2O Blk-3 8/25/99	<LOQ (0.0116)		NA	<LOQ (0.00625)		NA
	H2O Blk-4 8/25/99	<LOQ (0.0116)	<LOQ	NA	<LOQ (0.00625)	<LOQ	NA
Caspian Seal Blood	W2491-40,J 53	0.0180			<LOQ (0.00625)		
	W2491-41,J 11	<LOQ (0.0116)			<LOQ (0.00625)		
	W2491-42,J 46	<LOQ (0.0116)			<LOQ (0.00625)		
	W2491-43,J 17	0.0166			<LOQ (0.00625)		
	W2491-44,J 13	<LOQ (0.0116)			<LOQ (0.00625)		
	W2491-45,J 8	0.0131			<LOQ (0.00625)		
	W2491-46,J 12	<LOQ (0.0116)			<LOQ (0.00625)		
	W2491-47,J 9	<LOQ (0.0116)			<LOQ (0.00625)		
	W2491-48,J 14	<LOQ (0.0116)			<LOQ (0.00625)		
	W2491-49,J 18	<LOQ (0.0116)			<LOQ (0.00625)		
	W2491-50,J 52	<LOQ (0.0116)			<LOQ (0.00625)		
	W2491-51,J 55	<LOQ (0.0116)			<LOQ (0.00625)		
	W2491-52,J 10	<LOQ (0.0116)		NA	<LOQ (0.00625)		NA
	W2491-53,J 15	<LOQ (0.0116)	<LOQ - 3 Outliers	NA	<LOQ (0.00625)	<LOQ	NA

No curve analyzed for PFHS, PFDS. PFHS based on PFOS response.

No PFOS qualitative confirmation performed. Identifications are preliminary. Deviant Surrogate levels are not noted and were not confirmed.

Date Entered/By: 08/27/99, 09/01/99, 12/30/99, 01/20/00, 02/14/00 LAC
 Date Verified/By: 02/22/00 MEE

LOQ = Limit of Quantitation

NA = Not Applicable

RSD = Relative Standard Deviation

ND = Not Detected

D = Detected

PFOS = Perfluorooctanesulfonate

PFOSA = Perfluorooctanesulfonamide

POAA = Perfluorooctanoate

PFHS = Perfluorohexanesulfonate

Study:
 Product Number(Test Substance):
 Matrix:
 Method/Revision:
 Analytical Equipment System Number:
 Instrument Software/Version:
 Filename:
 R-Squared Value:
 Slope:
 Y-Intercept:
 Dates of Extraction/Analyst:
 Dates of Analysis/Analyst:
 Date of Data Reduction/Analyst:

GEN021 Various Matrices from MSU
 None
 Blood
 ETS-8-4.1 & ETS-8-5.1 using unextracted curves
 Amelia 062498
 Masslynx 3.2
 See list to right
 See Attachments
 See Attachments
 See Attachments
 08/25/99 MCH/KK/SAL/SEE
 08/26/99, 08/27/99, 08/28/99, 12/09/99 MEE/IAS
 08/27/99, 08/30/99, 08/31/99, 12/10/99, 01/20/00 MEE/IAS

Sample Data

BLOOD

Group Dose	Sample #	Concentration of POAA ug/mL	Mean POAA ug/mL	RSD Std. Dev.	Concentration of PFHS ug/mL	Mean PFHS ug/mL	RSD Std. Dev.
Method Blk	H2O Blk-1 8/25/99	0.00629			ND		
	H2O Blk-2 8/25/99	<LOQ (0.00599)			ND		
	H2O Blk-3 8/25/99	<LOQ (0.00599)		NA	ND		NA
	H2O Blk-4 8/25/99	<LOQ (0.00599)	<LOQ - 1 Outlier	NA	ND	ND	NA
Caspian Seal Blood	W2491-40,J 53	<LOQ (0.00599)			D		
	W2491-41,J 11	0.00759			D		
	W2491-42,J 46	<LOQ (0.00599)			ND		
	W2491-43,J 17	<LOQ (0.00599)			D		
	W2491-44,J 13	<LOQ (0.00599)			ND		
	W2491-45,J 8	<LOQ (0.00599)			ND		
	W2491-46,J 12	<LOQ (0.00599)			ND		
	W2491-47,J 9	<LOQ (0.00599)			ND		
	W2491-48,J 14	0.00728			ND		
	W2491-49,J 18	<LOQ (0.00599)			ND		
	W2491-50,J 52	0.0108			ND		
	W2491-51,J 55	0.0234			ND		
	W2491-52,J 10	<LOQ (0.00599)		NA	ND		NA
	W2491-53,J 15	<LOQ (0.00599)	<LOQ - 4 Outliers	NA	ND	ND - 3 Outliers	NA

No curve analyzed for PFHS, PFDS. PFHS based on PFOS response.

No PFOS qualitative confirmation performed. Identifications are preliminary.

Deviant Surrogate levels are not noted and were not confirmed.

Date Entered/By: 08/27/99, 09/01/99, 12/30/99, 01/20/00, 02/14/00 LAC

Date Verified/ By: 02/22/00 MEE

LOQ = Limit of Quantitation

NA = Not Applicable

RSD = Relative Standard Deviation

ND = Not Detected

D = Detected

PFOS = Perfluorooctanesulfonate

PFOSA = Perfluorooctanesulfonamide

POAA = Perfluorooctanoate

PFHS = Perfluorohexanesulfonate

FACT-GEN-021

Study:

Product Number(Test Substance):

Matrix:

Method/Revision:

Analytical Equipment System Number:

Instrument Software/Version:

Filename:

R-Squared Value:

Slope:

Y-Intercept:

Dates of Extraction/Analyst:

Dates of Analysis/Analyst:

Date of Data Reduction/Analyst:

Sample Data

GEN021 Various Matrices from MSU

None

Blood

ETS-B-4.1 & ETS-B-5.1 using unextracted curves

Amelia 062498

Mamlynx 3.2, 3.3

08/28/99, 12/09/99 IAS

08/28/99, 12/09/99 IAS

08/30/99, 12/10/99 IAS

See Attachments

08/25/99 MCH/KK/SAL/SEE

08/28/99, 12/09/99 IAS

08/30/99, 12/10/99 IAS

BLOOD

QC

Group Date	Sample #	Concentration of PFOS ug/mL or % Rec	Mean PFOS ug/mL or % Rec	RSD Std. Dev.	Concentration of PFOSA ug/mL or % Rec	Mean PFOSA ug/mL or % Rec	RSD Std. Dev.
Method Blk	H2O Blk-1 8/25/99	NA			NA		
	H2O Blk-2 8/25/99	NA			NA		
	H2O Blk-3 8/25/99	NA		NA	NA		NA
	H2O Blk-4 8/25/99	NA	NA	NA	NA	NA	NA
Caprine Seal Blood	W2491-40J 53-MS	164%			106%		
	W2491-41J 11-MS	115%			98%		
	W2491-42J 46-MS	101%			73%		
	W2491-43J 17-MS	2431%			1991%		
	W2491-44J 13-MS	67%			50%		
	W2491-45J 8-MS	103%			79%		
	W2491-46J 12-MS	130%			116%		
	W2491-47J 9-MS	103%			85%		
	W2491-48J 14-MS	103%			90%		
	W2491-49J 18-MS	73%			47%		
	W2491-50J 52-MS	65%			49%		
	W2491-51J 55-MS	17%			10%		
	W2491-52J 10-MS	12%	* outlier excluded	58%	10%	* outlier excluded	58%
	W2491-53J 15-MS	17%	84%	48%	12%	63%	37%

No curve analyzed for PFHS, PFDS. PFHS based on PFOS response.

No PFOS qualitative confirmation performed. Identifications are preliminary.

Deviant Surrogate levels are not noted and were not confirmed.

Date Entered/By: 02/16/00, 02/17/00 LAC

Date Verified/By: 02/22/00 MEE

LOQ = Limit of Quantitation

NA = Not Applicable

RSD = Relative Standard Deviation

ND = Not Detected

D = Detected

NS = Not Spiked

PFOS = Perfluorooctanesulfonate

PFOSA = Perfluorooctanesulfonamide

FOAA = Perfluorooctanoate

PFHS = Perfluorohexanesulfonate

FACT-GEN-021

Study:
 Product Number(Test Substance):
 Matrix:
 Method/Revision:
 Analytical Equipment System Number:
 Instrument Software/Version:
 Filename:
 R-Squared Value:
 Slope:
 Y-Intercept:
 Dates of Extraction/Analyst:
 Dates of Analysis/Analyst:
 Date of Data Reduction/Analyst:

GEN021 Various Matrices from MSU
 None
 Blood
 BTS-8-4.1 & BTS-8-5.1 using unextracted curves
 Arnelia 062498
 Masslynx 3.2, 3.3
 08/28/99, 12/09/99 IAS
 08/28/99, 12/09/99 IAS
 08/30/99, 12/10/99 IAS
 See Attachments
 08/25/99 MCH/KK/SAL/SEE
 08/28/99, 12/09/99 IAS
 08/30/99, 12/10/99 IAS

Sample Data

BLOOD

QC

Group Date	Sample #	Concentration of POAA ug/mL or % Rec	Mean POAA ug/mL or % Rec	RSD Std. Dev.	Concentration of PFHS ug/mL or % Rec	Mean PFHS ug/mL or % Rec	RSD Std. Dev.
Method Blk	H2O Blk-1 8/25/99	NA			NS		
	H2O Blk-2 8/25/99	NA			NS		
	H2O Blk-3 8/25/99	NA		NA	NS		NS
	H2O Blk-4 8/25/99	NA	NA	NA	NS	NS	NS
Caspian Seal Blood	W2491-40.J 53-MS	157%			NS		
	W2491-41.J 11-MS	113%			NS		
	W2491-42.J 46-MS	100%			NS		
	W2491-43.J 17-MS	2056%			NS		
	W2491-44.J 13-MS	89%			NS		
	W2491-45.J 8-MS	108%			NS		
	W2491-46.J 12-MS	174%			NS		
	W2491-47.J 9-MS	111%			NS		
	W2491-48.J 14-MS	112%			NS		
	W2491-49.J 18-MS	91%			NS		
	W2491-50.J 52-MS	78%			NS		
	W2491-51.J 55-MS	14%			NS		
	W2491-52.J 10-MS	19%	* outlier excluded	54%	NS		NS
	W2491-53.J 15-MS	19%	91%	50%	NS	NS	NS

No curve analyzed for PFHS, PFOS, PFOS based on PFOS response.
 No PFOS qualitative confirmation performed. Identifications are preliminary.
 Deviant Surrogate levels are not noted and were not confirmed.
 Date Entered/By: 02/16/00, 02/17/00 LAC
 Date Verified/By: 02/22/00 MEE

LOQ = Limit of Quantitation
 NA = Not Applicable
 RSD = Relative Standard Deviation
 ND = Not Detected
 D = Detected
 NS = Not Spiked

PFOS = Perfluorooctanesulfonate
 PFOSA = Perfluorooctanesulfonamide
 POAA = Perfluorooctanoate
 PFHS = Perfluorohexanesulfonate

Study:
 Product Number(Test Substance):
 Matrix:
 Method/Revision:
 Analytical Equipment System Number:
 Instrument Software/Version:
 Date of Extraction/Analyst:
 Date of Analysis/Analyst:
 Date of Data Reduction/Analyst:
 Sample Data

GEN021 Various Matrices from MSU
 None
 Various Matrices
 ETS-4-6.0 & ETS-8-7.0 using unextracted curves
 Amelia 062498
 Masslynx 3.2
 08/25/99 MCH/KEK/SAL/SEE
 08/26/99, 08/27/99, 08/28/99, 12/09/99 MEE/LAS
 08/27/99, 08/30/99, 08/31/99, 12/10/99, 01/20/00 MEE/LAS

LIVER/WHOLE BLOOD

Group Dose	Sample #	Concentration of PFOS ug/g or % Res.	Mean PFOS ug/g	RSD Std. Dev.	Concentration of PFOSA ug/g or % Res.	Mean PFOSA ug/g	RSD Std. Dev.
Method Blk	H2O Blk-1 8/25/99 H2O Blk-2 8/25/99 H2O Blk-3 8/25/99 H2O Blk-4 8/25/99	<LOQ (0.0348) <LOQ (0.0348) <LOQ (0.0348) <LOQ (0.0348)	<LOQ	<LOQ	<LOQ (0.0375) <LOQ (0.0375) <LOQ (0.0375) <LOQ (0.0375)	<LOQ	NA NA
California San Luis Liver	W2491-1,CSL 1448 W2491-4,CSL 1395 W2491-6,CSL 1020 W2491-7,CSL 2169 W2491-9,CSL 2839 W2491-10,CSL 2367	0.0384 0.0494 <LOQ (0.0348) <LOQ (0.0348) <LOQ (0.0348) <LOQ (0.0348)	<LOQ - 2 Outliers	NA 0.00773	0.0443 <LOQ (0.0375) <LOQ (0.0375) <LOQ (0.0375) <LOQ (0.0375) <LOQ (0.0375)	<LOQ - 1 Outlier	NA NA NA NA
Elephant Seal Liver	W2491-2,ES 1500 W2491-5,ES 1552 W2491-11,ES 808 W2491-12,ES 772 W2491-14,ES 782	<LOQ (0.0348) <LOQ (0.0348) <LOQ (0.0348) <LOQ (0.0348) <LOQ (0.0348)	<LOQ	NA NA	<LOQ (0.0375) <LOQ (0.0375) <LOQ (0.0375) <LOQ (0.0375) <LOQ (0.0375)	<LOQ	NA NA NA NA
Harbor Seal Liver	W2491-13,ES 1191 W2491-15,ES 1199	<LOQ (0.0348) <LOQ (0.0348)	<LOQ - 1 Outlier	NA NA	<LOQ (0.0375) <LOQ (0.0375)	<LOQ	NA NA
Coast Liver	W2491-1,NFS 100	0.133	0.133	NA	<LOQ (0.0375)	<LOQ	NA
Mink Liver	W2491-34,D1030 USFW3 W2491-35,D1146 USFW3 W2491-36,D1158 USFW3	4.85 2.41 0.587	2.62	81.7 2.14	<LOQ (0.0375) <LOQ (0.0375) <LOQ (0.0375)	<LOQ	NA NA NA
River Otter Liver	W2491-29,RAG 066 W2491-30,RAG 028 W2491-31,RAG 148 W2491-32,RAG 230 W2491-33,RAG 237	0.779 0.994 0.189 0.0334 0.151	0.329	116 0.382	0.0371 0.0448 0.0716 <LOQ (0.0375) 0.0393	0.0482 - 1 Outlier	33.1 0.0160
Sea Otter Liver	W2491-16,SO 12593-001 W2491-19,SO 11494-001 W2491-22,SO 11940-001 W2491-24,SO 11309-001 W2491-25,SO 12797-001 W2491-26,SO 13110-001 W2491-27,SO 12679-001 W2491-28,SO 12707-001	<LOQ (0.0348) <LOQ (0.0348) <LOQ (0.0348) <LOQ (0.0348) <LOQ (0.0348) <LOQ (0.0348) <LOQ (0.0348) <LOQ (0.0348)	<LOQ	NA NA	0.0806 <LOQ (0.0375) <LOQ (0.0375) <LOQ (0.0375) <LOQ (0.0375) <LOQ (0.0375) <LOQ (0.0375) <LOQ (0.0375)	<LOQ - 1 Outlier	NA NA NA NA
Turtle Liver	W2491-37, Male Turtle (-2.8), Turtle Liver W2491-38, Male Turtle (2.12), Turtle Liver W2491-39, Female Turtle (-3.9), Turtle Liver	0.239 0.358 0.099	0.232	56.0 0.130	<LOQ (0.0375) <LOQ (0.0375) <LOQ (0.0375)	<LOQ	NA NA NA
Sea Otter Brain	W2491-18,SO 12593-001, Sea Otter Brain W2491-21,SO 11494-001, Sea Otter Brain	<LOQ (0.0348) <LOQ (0.0348)	<LOQ	NA NA	0.0664 <LOQ (0.0375)	NA	NA NA
Sea Otter Kidney	W2491-17,SO 12593-001, Sea Otter Kidney W2491-20,SO 11494-001, Sea Otter Kidney W2491-23,SO 11940-001, Sea Otter Kidney	<LOQ (0.0348) <LOQ (0.0348) <LOQ (0.0348)	<LOQ	NA NA	<LOQ (0.0375) <LOQ (0.0375) <LOQ (0.0375)	<LOQ	NA NA NA
Whole Blood	W2491-54, Cormorant DCCO L. Chertsey W2491-55, Cormorant DCCO Hymn Island, Lake Sup W2491-56, Otter DCCO Great Lakes	0.190 0.0422 0.0392	0.0904	95.2 0.0661	<LOQ (0.0375) <LOQ (0.0375) 0.112	<LOQ - 1 Outlier	NA NA NA

PFOS/PFHS = no curve analyzed PFHS based on PFOS response.

No PFOS qualitative confirmation performed. Identifications are preliminary.
 Derivat Surrogate levels are not noted and were not confirmed.

Date Entered/Analyst: 08/17/99, 09/01/99, 12/09/99, 01/20/00, 02/14/00 LAC
 Date Verified/Analyst: 08/22/00 MEE

LOQ = Limit of Quantitation
 RSD = Relative Standard Deviation
 NA = Not Applicable
 ND = Not Detected
 D = Detected

PFOS = Perfluorooctanesulfonate
 PFOSA = Perfluorooctanesulfonamide
 FOAA = Perfluorooctanoate
 PFHS = Perfluorohexanesulfonate

Study:
Product Number(Test Substance):
Matrix:
Method/Revision:
Analytical Equipment System Number:
Instrument Software/Version:
Date of Extraction/Analyst:
Date of Analysis/Analyst:
Date of Data Reduction/Analyst:
Sample Data

GEN021 Various Matrices from MSU
None
Various Matrices
ETS-4-6.0 & ETS-4-7.0 using unextracted curves
Amelia 062498
Masslynx 3.2
06/25/99 MCH/KK/SAL/SEE
06/26/99, 08/27/99, 08/28/99, 12/09/99 MEB/IAS
08/27/99, 08/30/99, 08/31/99, 12/10/99, 01/20/00 MEB/IAS

LIVER/WHOLE BLOOD

Group Dose	Sample #	Concentration of POAA ug/g or % Res.	Mean POAA ug/g	RSD Std. Dev.	Concentration of PFHS ug/g or % Res.	Mean PFHS ug/g	RSD Std. Dev.
Method Btk	H2O BB-1 8/25/99 H2O BB-2 8/25/99 H2O BB-3 8/25/99 H2O BB-4 8/25/99	0.00602 <LOQ (0.0359) <LOQ (0.0359) <LOQ (0.0359)	<LOQ - 1 Outlier	NA NA	ND ND ND ND	ND	NA NA
California Sea Lion Liver	W2491-3,CSL 3448 W2491-4,CSL 3395 W2491-6,CSL 3020 W2491-7,CSL 2169 W2491-9,CSL 2839 W2491-10,CSL 2367	<LOQ (0.0359) 0.0409 <LOQ (0.0359) <LOQ (0.0359) <LOQ (0.0359) <LOQ (0.0359)	<LOQ - 1 Outlier	NA NA	ND ND ND ND D ND	ND - 1 Outlier	NA NA
Elephant Seal Liver	W2491-2,ES 1500 W2491-5,ES 1552 W2491-11,ES 806 W2491-12,ES 772 W2491-14,ES 782	<LOQ (0.0359) <LOQ (0.0359) <LOQ (0.0359) <LOQ (0.0359) <LOQ (0.0359)	<LOQ	NA NA	ND ND ND ND ND	ND	NA NA
Harbor Seal Liver	W2491-8,HS W2491-13,HS 1191 W2491-15,HS 1199	<LOQ (0.0359) <LOQ (0.0359) <LOQ (0.0359)	<LOQ	NA NA	ND ND ND	ND	NA NA
Coast Liver	W2491-13,PS 100	<LOQ (0.0359)	<LOQ	NA	D	D	NA
Mink Liver	W2491-34,D1000 USFWS W2491-35,D1146 USFWS W2491-36,D1158 USFWS	<LOQ (0.0359) <LOQ (0.0359) <LOQ (0.0359)	<LOQ	NA NA	D ND ND	ND - 1 Outlier	NA NA
River Otter Liver	W2491-29,RAG 066 W2491-30,RAG 028 W2491-31,RAG 148 W2491-32,RAG 230 W2491-33,RAG 237	<LOQ (0.0359) <LOQ (0.0359) <LOQ (0.0359) <LOQ (0.0359) <LOQ (0.0359)	<LOQ	NA NA	D D D ND D	D - 1 Outlier	NA NA
Sea Otter Liver	W2491-16,SO 12593-001 W2491-19,SO 11494-001 W2491-22,SO 11940-001 W2491-24,SO 11309-001 W2491-25,SO 12797-001 W2491-26,SO 13110-001 W2491-27,SO 12679-001 W2491-28,SO 12707-001	<LOQ (0.0359) <LOQ (0.0359) <LOQ (0.0359) <LOQ (0.0359) <LOQ (0.0359) <LOQ (0.0359) <LOQ (0.0359) <LOQ (0.0359)	<LOQ	NA NA	ND D ND ND D ND ND ND	ND - 2 Outliers	NA NA
Turtle Liver	W2491-37,Male Turtle (-2.3),Turtle Liver W2491-38,Male Turtle (2.12),Turtle Liver W2491-39,Female Turtle (-3.9),Turtle Liver	<LOQ (0.0359) <LOQ (0.0359) <LOQ (0.0359)	<LOQ	NA NA	ND D D	D - 1 Outlier	NA NA
Sea Otter Brain	W2491-18,SO 12593-001,Sea Otter Brain W2491-21,SO 11494-001,Sea Otter Brain	<LOQ (0.0359) <LOQ (0.0359)	<LOQ	NA NA	ND ND	ND	NA NA
Sea Otter Kidney	W2491-17,SO 12593-001,Sea Otter Kidney W2491-20,SO 11494-001,Sea Otter Kidney W2491-23,SO 11940-001,Sea Otter Kidney	<LOQ (0.0359) <LOQ (0.0359) <LOQ (0.0359)	<LOQ	NA NA	ND ND ND	ND	NA NA
Whale Blood	W2491-54,Commons DCCO L Charity W2491-55,Commons DCCO Hyatt Island, Lake Sup W2491-56,Outer DCCO Great Lakes	<LOQ (0.0359) <LOQ (0.0359) <LOQ (0.0359)	<LOQ	NA NA	ND D D	D - 1 Outlier	NA NA

PFOS/PFHS = no curve analyzed PFHS based on PFOS response.

No PFOS qualitative confirmation performed. Identifications are preliminary.

Deviant Surrogate levels are not noted and were not confirmed.

Date Entered/Analyst: 06/27/99, 09/01/99, 12/30/99, 01/20/00, 02/14/00 LAC
Date Verified/Analyst: 02/22/00 MEE

LOQ = Limit of Quantitation
RSD = Relative Standard Deviation
NA = Not Applicable
ND = Not Detected
D = Detected

PFOS = Perfluorooctanesulfonate
PFOSA = Perfluorooctanesulfonamide
POAA = Perfluorooctanoate
PFHS = Perfluorohexanesulfonate

Study:
 Product Number(Test Substance):
 Matrix:
 Method/Revision:
 Analytical Equipment System Number:
 Instrument Software/Version:
 Date of Extraction/Analyst:
 Date of Analysis/Analyst:
 Date of Data Reduction/Analyst:
Sample Data

GEN021 Various Matrices from MSU
 None
 Various Matrices
 ETS-8-6.0 & ETS-8-7.0 using unextracted curves
 Amelia 062498
 Masslynx 3.2, 3.3
 08/25/99 MCH/KK/SAL/SEE
 08/28/99, 12/09/99 LAS
 08/30/99, 12/10/99 LAS

LIVER/WHOLE BLOOD

QC

Group Dose	Sample #	Concentration of PFOS ug/g or % Rec.	Mean PFOS Recovery	RSD Std. Dev.	Concentration of PFOSA ug/g or % Rec.	Mean PFOSA ug/g	RSD Std. Dev.
Method Blk	H2O Blk-1 8/25/99 H2O Blk-2 8/25/99 H2O Blk-3 8/25/99 H2O Blk-4 8/25/99	NA NA NA NA	NA	NA NA	NA NA NA NA	NA	NA NA
California Sea Lion Liver	W2491-3,CSL 3448-MS W2491-4,CSL 3395-MS W2491-6,CSL 3020-MS W2491-7,CSL 2169-MS W2491-9,CSL 2839-MS W2491-10,CSL 2367-MS	91% 78% 69% 45% 31% 64%	63%	35% 22%	66% 52% 48% 30% 15% 43%	42%	42% 18%
Elephant Seal Liver	W2491-2,ES 1500-MS W2491-5,ES 1552-MS W2491-11,ES 808-MS W2491-12,ES 772-MS W2491-14,ES 782-MS	105% 77% 51% 44% 33%	62%	47% 29%	78% 65% 35% 18% 29%	45%	56% 25%
Harbor Seal Liver	W2491-4,HS-MS W2491-13,HS 1191-MS W2491-15,HS 1199-MS	35% 81% 50%	55%	42% 23%	42% 79% 42%	54%	39% 21%
Goat Liver	W2491-1,NFS 100-MS	57%	NA	NA	63%	NA	NA
Mink Liver	W2491-34,D1030 USFWS-MS W2491-35,D1146 USFWS-MS W2491-36,D1158 USFWS-MS	NR NR NR	NR	NA NA	NR NR NR	NR	NA NA
River Otter Liver	W2491-29,RAG 066-MS W2491-30,RAG 028-MS W2491-31,RAG 148-MS W2491-32,RAG 230-MS W2491-33,RAG 237-MS	NR NR 48% 38% 37%	41%	15% 6%	NR NR 47% 34% 29%	36%	26% 9%
Sea Otter Liver	W2491-16,SO 12593-001-MS W2491-19,SO 11494-001-MS W2491-22,SO 11940-001-MS W2491-24,SO 11309-001-MS W2491-25,SO 12797-001-MS W2491-26,SO 13110-001-MS W2491-27,SO 12679-001-MS W2491-28,SO 12707-001-MS	42% 61% 44% 27% 34% 69% 36% 65%	47%	33% 16%	11% 50% 35% 16% 22% 44% 26% 33%	30%	45% 13%
Turtle Liver	W2491-37,Male Turtle (2.8)-MS W2491-38,Male Turtle (2.12)-MS W2491-39,Female Turtle (-3.9)-MS	37% 35% 51%	41%	20% 8%	42% 51% 39%	44%	15% 6%
Sea Otter Brain	W2491-18,SO 12593-001-MS W2491-21,SO 11494-001-MS	42% 46%	44%	7% 3%	-3% 27%	12%	171% 21%
Sea Otter Kidney	W2491-17,SO 12593-001-MS W2491-20,SO 11494-001-MS W2491-23,SO 11940-001-MS	61% 75% 24%	53%	49% 26%	66% 57% 14%	46%	61% 28%
Whole Blood	W2491-54,Cormorant DCCO I Charity-MS W2491-55,Cormorant DCCO Hym Island, Lake Sup-MS W2491-56,Otter DCCO Great Lakes-MS	63% 69% 115%	82%	34% 28%	65% 64% 104%	77%	29% 23%

PFDS/FFHS = no curve analyzed, FFHS based on PFOS response.

No PFOS qualitative confirmation performed. Identifications are preliminary.

NR = Not reported, appears the spike wasn't detectable from endogenous levels.

LOQ = Limit of Quantitation

RSD = Relative Standard Deviation

NA = Not Applicable

NS = Not spiked

PFOS = Perfluorooctanesulfonate

PFOSA = Perfluorooctanesulfonamide

POAA = Perfluorooctanoate

FFHS = Perfluorohexanesulfonate

Date Entered/Analyst:

02/16/00, 02/17/00 LAC

Date Verified/Analyst:

20/02/00 MEE

Study:
 Product Number(Test Substance):
 Matrix:
 Method/Revision:
 Analytical Equipment System Number:
 Instrument Software/Version:
 Date of Extraction/Analyst:
 Date of Analysis/Analyst:
 Date of Data Reduction/Analyst:
 Sample Data

GEN021 Various Matrices from MSU
 None
 Various Matrices
 ETS-8-6.0 & ETS-8-7.0 using unextracted curves
 Amelia 062498
 Masslynx 3.2, 3.3
 08/28/99 MCH/KK/SAL/SEE
 08/28/99, 12/09/99 IAS
 08/30/99, 12/10/99 IAS

LIVER/WHOLE BLOOD

QC

Group Dose	Sample #	Concentration of POAA ug/g or % Rec.	Mean POAA ug/g	RSD Std. Dev.	Concentration of PFHS ug/g or % Rec.	Mean PFHS ug/g	RSD Std. Dev.
Method Blk	H2O Blk-1 8/25/99 H2O Blk-2 8/25/99 H2O Blk-3 8/25/99 H2O Blk-4 8/25/99	NA NA NA NA	NA	NA NA	NS NS NS NS	NS	NA NA
California Sea Lion Liver	W2491-3,CSL 3448-MS W2491-4,CSL 3395-MS W2491-6,CSL 3020-MS W2491-7,CSL 2169-MS W2491-9,CSL 2839-MS W2491-10,CSL 2367-MS	109% 85% 68% 45% 20% 64%		47% 31%	NS NS NS NS NS NS	NS	NA NA
Elephant Seal Liver	W2491-2,ES 1500-MS W2491-5,ES 1552-MS W2491-11,ES 308-MS W2491-12,ES 772-MS W2491-14,ES 782-MS	100% 80% 44% 31% 36%		52% 30%	NS NS NS NS NS	NS	NA NA
Harbor Seal Liver	W2491-4,HS-MS W2491-13,HS 1191-MS W2491-15,HS 1199-MS	48% 94% 58%		37% 25%	NS NS NS	NS	NA NA NA
Coast Liver	W2491-1,NFS 100-MS	91%	NA	NA	NS	NS	NA
Mink Liver	W2491-34,D1030 USFWS-MS W2491-35,D1146 USFWS-MS W2491-36,D1158 USFWS-MS	NR NR NR	NA NA	NA NA	NS NS NS	NS	NA NA NA
River Otter Liver	W2491-29,RAG 066-MS W2491-30,RAG 028-MS W2491-31,RAG 148-MS W2491-32,RAG 230-MS W2491-33,RAG 237-MS	21% 46% 69% 52% 70%		39% 20%	NS NS NS NS NS	NS	NA NA
Sea Otter Liver	W2491-16,SO 12593-001-MS W2491-19,SO 11494-001-MS W2491-22,SO 11940-001-MS W2491-24,SO 11309-001-MS W2491-25,SO 12797-001-MS W2491-26,SO 13110-001-MS W2491-27,SO 12679-001-MS W2491-28,SO 12707-001-MS	47% 64% 47% 22% 30% 83% 44% 64%		39% 20%	NS NS NS NS NS NS NS NS	NS	NA NA
Turtle Liver	W2491-37,Male Turtle (-2,8)-MS W2491-38,Male Turtle (2,12)-MS W2491-39,Female Turtle (-3,9)-MS	48% 71% 42%		29% 15%	NS NS NS	NS	NA NA NA
Sea Otter Brain	W2491-18,SO 12593-001-MS W2491-21,SO 11494-001-MS	41% 50%	46%	13% 6%	NS NS	NS	NA NA
Sea Otter Kidney	W2491-17,SO 12593-001-MS W2491-20,SO 11494-001-MS W2491-23,SO 11940-001-MS	94% 95% 9%	66%	75% 49%	NS NS NS	NS	NA NA NA
Whole Blood	W2491-54,Cormorant DCCO L Charity-MS W2491-55,Cormorant DCCO Hymn Island, Lake Sup-MS W2491-56,Otter DCCO Great Lakes-MS	118% 122% 123%		2% 3%	NS NS NS	NS	NA NA NA

PFDS/PFHS = no curve analyzed, PFHS based on PFOS response.

No PFOS qualitative confirmation performed. Identifications are preliminary.

NR = Not reported, appears the spike wasn't detectable from endogenous levels.

LOQ = Limit of Quantitation

RSD = Relative Standard Deviation

NA = Not Applicable

NS = Not spiked

PFOS = Perfluorooctanesulfonate

PFOSA = Perfluorooctanesulfonamide

POAA = Perfluorooctanoate

PFHS = Perfluorohexanesulfonate

Date Entered/Analyst:

02/16/00, 02/17/00 LAC

Date Verified/Analyst:

20/22/00 MEE

C

Study:
 Product Number(Test Substance):
 Matrix:
 Method/Revision:
 Analytical Equipment System Number:
 Instrument Software/Version:
 Filename:
 R-Squared Value:
 Slope:
 Y-Intercept:
 Dates of Extraction/Analyst:
 Dates of Analysis/Analyst:
 Date of Data Reduction/Analyst:

GEN024 MSU Environmental Samples
 Various
 Various
 ETS-4-4.1 & ETS-4-5.1 using unextracted curves
 Amelia 061496
 Masslynx 3.3
 See Attachments
 See Attachments
 See Attachments
 See Attachments
 10/12/99 SAL/KK
 10/15/99, 10/19/99 IAS/MMH, 12/13/99 IAS
 10/18/99, 10/20/99 HOI, 12/16/99 MMH

Sample Data

SERA

Group Dose	Sample #	Concentration of PFOS ug/mL or % Rec	Mean PFOS ug/mL	RSD Std. Dev. RPD-MS/MSD	Concentration of PFOS ug/mL or % Rec	Mean PFOS ug/mL	RSD Std. Dev. RPD-MS/MSD
Method Blk	Bird10129-WBlt-5-1	<LOQ (0.00114 ug/mL)	<LOQ	<LOQ	<LOQ (0.00114 ug/mL)	<LOQ	<LOQ
Matrix Blk	NE	NE	NE	NE	NE	NE	NE
MS/MSD 250 ppb	Bird 040-Alb sera-MS-250 ppb-5-1	81%	89%	17%	81%	75%	17%
	Bird 040-Alb sera-MSD-250 ppb-5-1	96%			59%		
	Bird 060-E plasma-MS-250 ppb-5-2	11%	17%	69%	63%	62%	11%
	Bird 060-E plasma-MSD-250 ppb-5-2	23%			76%		
	Bird 054-C blood-MS-250 ppb-5	84%	19%	678%	14%	45%	139%
	Bird 054-C blood-MSD-250 ppb-5	46%					
Albatross	Bird 034-Albatross Chick sera	0.0393			<LOQ (0.00114 ug/mL)		
	Bird 035-Albatross plasma	0.0256			<LOQ (0.00114 ug/mL)		
	Bird 036-Albatross sera	0.00755			<LOQ (0.00114 ug/mL)		
	Bird 037-Albatross plasma	0.0196			<LOQ (0.00114 ug/mL)		
	Bird 038-Albatross sera	0.00348			<LOQ (0.00114 ug/mL)		
	Bird 039-Albatross sera	0.00677			<LOQ (0.00114 ug/mL)		
	Bird 040-Albatross sera	0.00621			<LOQ (0.00114 ug/mL)		
	Bird 041-Albatross sera	0.00430	Plasma	47.0	<LOQ (0.00114 ug/mL)	Plasma	NA
	Bird 042-Albatross sera	0.00402	0.0180	0.00847	<LOQ (0.00114 ug/mL)	<LOQ	NA
	Bird 043-Albatross sera	0.00546			<LOQ (0.00114 ug/mL)		
	Bird 044-Albatross sera	0.00897			<LOQ (0.00114 ug/mL)		
	Bird 045-Albatross sera	0.00657	Sera	115	<LOQ (0.00114 ug/mL)	Sera	NA
	Bird 046-Albatross plasma	0.00889	0.00926	0.0107	<LOQ (0.00114 ug/mL)	<LOQ	NA
Comorant	Bird 047-Comorant plasma	0.431			<LOQ (0.00114 ug/mL)		
	Bird 048-Comorant plasma	0.242		61.8	<LOQ (0.00114 ug/mL)		NA
	Bird 049-Comorant plasma	0.110	0.261	0.161	<LOQ (0.00114 ug/mL)	<LOQ	NA
	Bird 050-Comorant blood	0.0725			<LOQ (0.00114 ug/mL)		
	Bird 051-Comorant blood	0.218			<LOQ (0.00114 ug/mL)		
	Bird 052-Comorant blood	0.153			<LOQ (0.00114 ug/mL)		
	Bird 053-Comorant blood	<LOQ (0.00116 ug/mL)			<LOQ (0.00114 ug/mL)		
	Bird 054-Comorant blood	0.144		44.5	<LOQ (0.00114 ug/mL)		NA
	Bird 055-Comorant blood	0.273	0.172 - 1 Outlier	0.0764	<LOQ (0.00114 ug/mL)	<LOQ	NA
Herring	Bird 056-Herring Gull plasma	0.453		34.1	0.00243		NA
	Bird 057-Herring Gull plasma	0.277	0.365	0.125	<LOQ (0.00114 ug/mL)	NA	NA
	Bird 058-Herring Gull blood	0.0790		12.2	<LOQ (0.00114 ug/mL)		NA
	Bird 059-Herring Gull blood	0.0664	0.0727	0.00888	<LOQ (0.00114 ug/mL)	<LOQ	NA
Bald Eagle	Bird 060-Bald Eagle plasma	0.449			<LOQ (0.00114 ug/mL)		
	Bird 061-Bald Eagle plasma	0.316			<LOQ (0.00114 ug/mL)		
	Bird 062-Bald Eagle plasma	0.185			0.00148		
	Bird 063-Bald Eagle plasma	0.167			<LOQ (0.00114 ug/mL)		
	Bird 064-Bald Eagle plasma	0.0402			<LOQ (0.00114 ug/mL)		
	Bird 065-Bald Eagle plasma	0.0147			<LOQ (0.00114 ug/mL)		
	Bird 066-Bald Eagle plasma	0.122			<LOQ (0.00114 ug/mL)		
	Bird 067-Bald Eagle plasma	0.0865			<LOQ (0.00114 ug/mL)		
	Bird 068-Bald Eagle plasma	2.57			<LOQ (0.00114 ug/mL)		
	Bird 069-Bald Eagle plasma	2.35			<LOQ (0.00114 ug/mL)		
	Bird 070-Bald Eagle plasma	0.0324			<LOQ (0.00114 ug/mL)		
	Bird 071-Bald Eagle plasma	0.0908			<LOQ (0.00114 ug/mL)		
	Bird 072-Bald Eagle plasma	0.217			<LOQ (0.00114 ug/mL)		
	Bird 073-Bald Eagle plasma	0.168			0.00131		
	Bird 074-Bald Eagle plasma	0.163			<LOQ (0.00114 ug/mL)		
	Bird 075-Bald Eagle plasma	0.0822			<LOQ (0.00114 ug/mL)		
	Bird 076-Bald Eagle plasma	0.0238			<LOQ (0.00114 ug/mL)		
	Bird 077-Bald Eagle plasma	0.0573			<LOQ (0.00114 ug/mL)		
	Bird 078-Bald Eagle plasma	<LOQ (0.00116 ug/mL)			<LOQ (0.00114 ug/mL)		
	Bird 079-Bald Eagle plasma	0.286			<LOQ (0.00114 ug/mL)		
	Bird 080-Bald Eagle plasma	0.281			<LOQ (0.00114 ug/mL)		
	Bird 081-Bald Eagle plasma	0.649			0.00617		
	Bird 082-Bald Eagle plasma	0.335			<LOQ (0.00114 ug/mL)		
	Bird 083-Bald Eagle plasma	0.0718			<LOQ (0.00114 ug/mL)		
	Bird 084-Bald Eagle plasma	0.0381		183	<LOQ (0.00114 ug/mL)		NA
	Bird 085-Bald Eagle plasma	0.118	0.356 - 1 Outlier	0.651	<LOQ (0.00114 ug/mL)	<LOQ - 3 Outliers	NA

No PFOS qualitative confirmation performed. Identifications are preliminary.
 * Appears to not have been spiked. LAC 10/19/99

Date Entered/By: 10/18/99, 10/20/99 LAC 12/30/99 MMH
 Date Verified/ By: 2/17/00 mmh

NE = Not extracted
 NA = Not Applicable
 NS = Not Spiked
 LOQ = Limit of Quantitation
 RSD = Relative Standard Deviation
 RPD = Relative Percent Difference

PFOS = Perfluorooctanesulfonate
 PFHS = Perfluorohexanesulfonate
 FOAA = Perfluorooctanoate
 PFOA = Perfluorooctanesulfonamide

Study:
 Product Number(Test Substance):
 Matrix:
 Method/Revision:
 Analytical Equipment System Number:
 Instrument Software/Version:
 Filename:
 R-Squared Value:
 Slope:
 Y-Intercept:
 Dates of Extraction/Analyst:
 Dates of Analysis/Analyst:
 Date of Data Reduction/Analyst:

GEN024 MSU Environmental Samples
 Various
 Various
 ETS-4-4.1 & ETS-8-5.1 using unextracted curves
 Amelia 062498
 Masslynx 3.3
 See Attachments
 See Attachments
 See Attachments
 See Attachments
 10/12/99 SAL/KK
 10/15/99, 10/19/99 IAS/MMH, 12/13/99 IAS
 10/18/99, 10/20/99 HOJ, 12/14/99 MMH

Sample Data

SERA

Group Dose	Sample #	Concentration of POAA ug/mL or % Rec	Mean POAA ug/mL	RSD Std. Dev. RPD-MS/MSD	Concentration of PFOSA ug/mL or % Rec	Mean PFOSA ug/mL	RSD Std. Dev. RPD-MS/MSD
Method Blk	Bird10129-WBIL-5-1	<LOQ (0.0299 ug/mL)	<LOQ	<LOQ	<LOQ (0.00625 ug/mL)	<LOQ	<LOQ
Matrix Blk	NE	NE	NE	NE	NE	NE	NE
MS/MSD 250 ppb	Bird 040-Alb sera-MS-250 ppb-5-1	89%			70%		
	Bird 040-Alb sera-MSD-250 ppb-5-1	105%	97%	17%	80%	75%	14%
	Bird 060-E plasma-MS-250 ppb-5-2	77%			59%		
	Bird 060-E plasma-MSD-250 ppb-5-2	84%	80%	9%	65%	62%	9%
	Bird 054-C blood-MS-250 ppb-5	90%			77%		
	Bird 054-C blood-MSD-250 ppb-5	23%	56%	119%	14%	45%	140%
Albatross	Bird 034-Albatross Chick sera	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		
	Bird 035-Albatross plasma	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		
	Bird 036-Albatross sera	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		
	Bird 037-Albatross plasma	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		
	Bird 038-Albatross sera	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		
	Bird 039-Albatross sera	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		
	Bird 040-Albatross sera	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		
	Bird 041-Albatross sera	<LOQ (0.0299 ug/mL)	Plasma	NA	<LOQ (0.00625 ug/mL)	Plasma	NA
	Bird 042-Albatross sera	<LOQ (0.0299 ug/mL)	<LOQ	NA	<LOQ (0.00625 ug/mL)	<LOQ	NA
	Bird 043-Albatross sera	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		
	Bird 044-Albatross sera	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		
	Bird 045-Albatross sera	<LOQ (0.0299 ug/mL)	Sera	NA	<LOQ (0.00625 ug/mL)	Sera	NA
	Bird 046-Albatross plasma	<LOQ (0.0299 ug/mL)	<LOQ	NA	<LOQ (0.00625 ug/mL)	<LOQ	NA
Comorant	Bird 047-Comorant plasma	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		
	Bird 048-Comorant plasma	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		
	Bird 049-Comorant plasma	<LOQ (0.0299 ug/mL)	<LOQ	NA	<LOQ (0.00625 ug/mL)	<LOQ	NA
	Bird 050-Comorant blood	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		
	Bird 051-Comorant blood	<LOQ (0.0299 ug/mL)			0.0145		
	Bird 052-Comorant blood	<LOQ (0.0299 ug/mL)			0.00871		
	Bird 053-Comorant blood	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		
	Bird 054-Comorant blood	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		82.7
	Bird 055-Comorant blood	0.0489	<LOQ - 1 Outlier	NA	0.0426	0.0219 - 3 Outliers	0.0181
Herring	Bird 056-Herring Gull plasma	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		
	Bird 057-Herring Gull plasma	<LOQ (0.0299 ug/mL)	<LOQ	NA	<LOQ (0.00625 ug/mL)	<LOQ	NA
	Bird 058-Herring Gull blood	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		
	Bird 059-Herring Gull blood	<LOQ (0.0299 ug/mL)	<LOQ	NA	<LOQ (0.00625 ug/mL)	<LOQ	NA
Bald Eagle	Bird 060-Bald Eagle plasma	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		
	Bird 061-Bald Eagle plasma	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		
	Bird 062-Bald Eagle plasma	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		
	Bird 063-Bald Eagle plasma	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		
	Bird 064-Bald Eagle plasma	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		
	Bird 065-Bald Eagle plasma	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		
	Bird 066-Bald Eagle plasma	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		
	Bird 067-Bald Eagle plasma	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		
	Bird 068-Bald Eagle plasma	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		
	Bird 069-Bald Eagle plasma	<LOQ (0.0299 ug/mL)			0.0751		
	Bird 070-Bald Eagle plasma	<LOQ (0.0299 ug/mL)			0.0996		
	Bird 071-Bald Eagle plasma	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		
	Bird 072-Bald Eagle plasma	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		
	Bird 073-Bald Eagle plasma	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		
	Bird 074-Bald Eagle plasma	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		
	Bird 075-Bald Eagle plasma	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		
	Bird 076-Bald Eagle plasma	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		
	Bird 077-Bald Eagle plasma	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		
	Bird 078-Bald Eagle plasma	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		
	Bird 079-Bald Eagle plasma	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		
	Bird 080-Bald Eagle plasma	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		
	Bird 081-Bald Eagle plasma	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		
	Bird 082-Bald Eagle plasma	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		
	Bird 083-Bald Eagle plasma	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		
	Bird 084-Bald Eagle plasma	<LOQ (0.0299 ug/mL)			<LOQ (0.00625 ug/mL)		
	Bird 085-Bald Eagle plasma	<LOQ (0.0299 ug/mL)	<LOQ	NA	<LOQ (0.00625 ug/mL)	<LOQ - 2 Outliers	NA
				NA			NA

No PFOS qualitative confirmation performed. Identifications are preliminary.

* Appears to not have been spiked. LAC 10/19/99

Date Entered/By: 10/18/99, 10/20/99 LAC 12/30/99 MMH
 Date Verified/By: 2/17/00 mmh

NE = Not extracted
 NA = Not Applicable
 NS = Not Spiked
 LOQ = Limit of Quantitation
 RSD = Relative Standard Deviation
 RPD = Relative Percent Difference

PFOS = Perfluorooctanesulfonate
 PFHS = Perfluorohexanesulfonate
 POAA = Perfluorooctanoate
 PFOSA = Perfluorooctanesulfonamide

D



Wilson Jones . *Quick Reference Index System*

© 1991 Wilson Jones Company

3M_MN01666022

2814.0026

Study:
Product Number/Test Substance):
Matrix:
Method/Revision:
Analytical Equipment System Number:
Instrument Software/Version:
Date of Extraction/Analyst:
Date of Analysis/Analyst:
Date of Data Reduction/Analyst:

GEN024 MSU Environmental Samples
Various
Various
ETS-8-6.0 & ETS-8-7.0 using unextracted curves
Ametek 062496
Masslynx 3.3
10/12/99 SAL/KK
10/14/99, 12/13/99 HOJ/LAS
10/15/99, 11/15/99, 12/14/99 HOJ/MMH

Sample Data

LIVER, KIDNEY, YOLK

Group Dose	Sample #	Concentration of PFOS ug/g or % Rec.	Mean PFOS ug/g	RSD Std. Dev. MS/MSD RPD	Concentration of PFHS ug/g or % Rec.	Mean PFHS ug/g	RSD Std. Dev. MS/MSD RPD
Method Blk	Bird10129-wbtk-5-1 Bird10129-wbtk-6-1	<LOQ (0.0348 ug/g) <LOQ (0.0348 ug/g)	<LOQ	NA	<LOQ (0.0683 ug/g) <LOQ (0.0683 ug/g)	<LOQ	NA
QC 250 ng/g	Bird001-Loon Lvr-MS	28%	28%	3%	54%	60%	11%
	Bird001-Loon Lvr-MSD	28%	28%	3%	65%	70%	20%
	Bird023-Albatross Kidney-MS	71%	71%	20%	63%	73%	5%
	Bird023-Albatross Kidney-MSD	87%	79%	20%	77%	73%	5%
	Bird030-Cormorant Yolk-MS	121%	128%	10%	71%	73%	5%
Liver	Bird001-Loon	0.345			<LOQ (0.0683 ug/g)		
	Bird002-Loon	0.689			<LOQ (0.0683 ug/g)		
	Bird003-Loon	0.185			<LOQ (0.0683 ug/g)		
	Bird004-Loon	0.199			<LOQ (0.0683 ug/g)		
	Bird005-Loon	0.202			<LOQ (0.0683 ug/g)		
	Bird006-Loon	0.105			<LOQ (0.0683 ug/g)		
	Bird007-Loon	<LOQ (0.0348 ug/g)	0.287 - 2 Outliers	73.6	<LOQ (0.0683 ug/g)	<LOQ	NA
	Bird008-Loon	<LOQ (0.0348 ug/g)		0.212	<LOQ (0.0683 ug/g)	<LOQ	NA
Liver	Bird009-Brown Pelican	0.0460		103	<LOQ (0.0683 ug/g)		
	Bird010-Brown Pelican	0.294	0.170	0.175	<LOQ (0.0683 ug/g)	<LOQ	NA
Liver	Bird011-Albatross	0.617			<LOQ (0.0683 ug/g)		
	Bird012-Albatross	<LOQ (0.0348 ug/g)			<LOQ (0.0683 ug/g)		
	Bird013-Albatross	<LOQ (0.0348 ug/g)			<LOQ (0.0683 ug/g)		
	Bird014-Albatross	<LOQ (0.0348 ug/g)			<LOQ (0.0683 ug/g)		
	Bird015-Albatross	<LOQ (0.0348 ug/g)			<LOQ (0.0683 ug/g)		
	Bird016-Albatross	<LOQ (0.0348 ug/g)			<LOQ (0.0683 ug/g)		
	Bird020-Albatross	<LOQ (0.0348 ug/g)			<LOQ (0.0683 ug/g)		
	Bird021-Albatross	<LOQ (0.0348 ug/g)	<LOQ - 1 Outlier	NA	<LOQ (0.0683 ug/g)	<LOQ	NA
Kidney	Bird022-Albatross	<LOQ (0.0348 ug/g)			<LOQ (0.0683 ug/g)		
	Bird017-Albatross	<LOQ (0.0348 ug/g)			<LOQ (0.0683 ug/g)		
	Bird018-Albatross	<LOQ (0.0348 ug/g)			<LOQ (0.0683 ug/g)		
	Bird019-Albatross	<LOQ (0.0348 ug/g)			<LOQ (0.0683 ug/g)		
	Bird023-Albatross	<LOQ (0.0348 ug/g)			<LOQ (0.0683 ug/g)		
	Bird024-Albatross	<LOQ (0.0348 ug/g)			<LOQ (0.0683 ug/g)		
	Bird025-Albatross	<LOQ (0.0348 ug/g)			<LOQ (0.0683 ug/g)		
	Bird026-Albatross	<LOQ (0.0348 ug/g)	<LOQ	NA	<LOQ (0.0683 ug/g)	<LOQ	NA
Yolk	Bird027-Cormorant	<LOQ (0.0348 ug/g)			<LOQ (0.0683 ug/g)		
	Bird028-Cormorant	0.134		39.5	<LOQ (0.0683 ug/g)		
	Bird029-Cormorant	0.317	0.235 - 1 Outlier	0.0930	<LOQ (0.0683 ug/g)	<LOQ	NA
Yolk	Bird030-Cormorant	0.254			<LOQ (0.0683 ug/g)		
	Bird031-Gull	0.146		64.8	<LOQ (0.0683 ug/g)		
	Bird032-Gull	<LOQ (0.0348 ug/g)	0.0999 - 1 Outlier	0.0648	<LOQ (0.0683 ug/g)	<LOQ	NA
	Bird033-Gull	0.0541			<LOQ (0.0683 ug/g)		

No PFOS qualitative confirmation performed. Identifications are preliminary.

LOQ = Limit of Quantitation

RSD = Relative Standard Deviation

RPD = Relative Percent Difference

Date Entered/Analyst: 10/23/99 LAC, 12/30/99 MMH

Date Verified/Analyst: 2/17/00 MMH

NA = Not Applicable

PFOS = Perfluorooctanesulfonate

PFHS = Perfluorohexanesulfonate

FOAA = Perfluorooctanoate

PFOSA = Perfluorooctane sulfonamide

Study:
 Product Number(Test Substance):
 Matrix:
 Method/Revision:
 Analytical Equipment System Number:
 Instrument Software/Version:
 Date of Extraction/Analyst:
 Date of Analysis/Analyst:
 Date of Data Reduction/Analyst:
 Sample Data

GEN024 MSU Environmental Samples
 Various
 Various
 ETS-8-6.0 & ETS-8-7.0 using unextracted curves
 Amelia 062498
 Masslynx 3.3
 10/12/99 SAL/KK
 10/14/99, 12/13/99 HOI/IAS
 10/15/99, 11/15/99, 12/14/99 HOI/MMH

LIVER, KIDNEY, YOLK

Group Dose	Sample #	Concentration of POAA ug/g or % Rec.	Mean POAA ug/g	RSD Std. Dev. MS/MSD RPD	Concentration of PFOSA ug/g or % Rec.	Mean PFOSA ug/g	RSD Std. Dev. MS/MSD RPD
Method Blk	Bird10129-wbtk-5-1 Bird10129-wbtk-6-1	<LOQ (0.180 ug/g) <LOQ (0.180 ug/g)	<LOQ	NA	<LOQ (0.00750 ug/g) <LOQ (0.00750 ug/g)	<LOQ	NA
QC 250 ng/g	Bird001-Loon Lvr-MSD Bird001-Loon Lvr-MSD Bird023-Albatross Kidney-MSD Bird023-Albatross Kidney-MSD Bird030-Comorant Yolk-MSD Bird030-Comorant Yolk-MSD	101% 108% 113% 121% 105% 130%	104% 117% 117% 117% 105% 117%	6% 7% 7% 7% 62% 22%	53% 51% 56% 58% 62% 72%	52% 57% 57% 57% 67% 67%	4% 4% 5% 5% 15% 15%
Liver	Bird001-Loon Bird002-Loon Bird003-Loon Bird004-Loon Bird005-Loon Bird006-Loon Bird007-Loon Bird008-Loon	<LOQ (0.180 ug/g) <LOQ (0.180 ug/g) <LOQ (0.180 ug/g) <LOQ (0.180 ug/g) <LOQ (0.180 ug/g) <LOQ (0.180 ug/g) <LOQ (0.180 ug/g) <LOQ (0.180 ug/g)	<LOQ	NA NA NA NA NA NA NA NA	0.0153 <LOQ (0.00750 ug/g) 0.0147 0.0262 0.0213 0.0242 <LOQ (0.00750 ug/g) <LOQ (0.00750 ug/g)	0.0204 - 3 Outliers	25.5 0.00520
Liver	Bird009-Brown Pelican Bird010-Brown Pelican	<LOQ (0.180 ug/g) <LOQ (0.180 ug/g)	<LOQ	NA NA	<LOQ (0.00750 ug/g) 0.178	0.178 - 1 Outlier	NA
Liver	Bird011-Albatross Bird012-Albatross Bird013-Albatross Bird014-Albatross Bird015-Albatross Bird016-Albatross Bird020-Albatross Bird021-Albatross Bird022-Albatross	<LOQ (0.180 ug/g) <LOQ (0.180 ug/g) <LOQ (0.180 ug/g) <LOQ (0.180 ug/g) <LOQ (0.180 ug/g) <LOQ (0.180 ug/g) <LOQ (0.180 ug/g) <LOQ (0.180 ug/g) 0.182	<LOQ - 1 Outlier	NA NA NA NA NA NA NA NA NA	0.527 <LOQ (0.00750 ug/g) <LOQ (0.00750 ug/g) <LOQ (0.00750 ug/g) <LOQ (0.00750 ug/g) <LOQ (0.00750 ug/g) <LOQ (0.00750 ug/g) <LOQ (0.00750 ug/g) <LOQ (0.00750 ug/g) <LOQ (0.00750 ug/g)	<LOQ	NA
Kidney	Bird017-Albatross Bird018-Albatross Bird019-Albatross Bird023-Albatross Bird024-Albatross Bird025-Albatross Bird026-Albatross	<LOQ (0.180 ug/g) <LOQ (0.180 ug/g) <LOQ (0.180 ug/g) <LOQ (0.180 ug/g) <LOQ (0.180 ug/g) <LOQ (0.180 ug/g) <LOQ (0.180 ug/g)	<LOQ	NA NA NA NA NA NA NA	<LOQ (0.00750 ug/g) <LOQ (0.00750 ug/g) <LOQ (0.00750 ug/g) <LOQ (0.00750 ug/g) <LOQ (0.00750 ug/g) <LOQ (0.00750 ug/g) <LOQ (0.00750 ug/g)	<LOQ	NA
Yolk	Bird027-Comorant Bird028-Comorant Bird029-Comorant Bird030-Comorant	<LOQ (0.180 ug/g) 0.245 <LOQ (0.180 ug/g) 0.192	0.218 - 2 outliers	17.2 0.0374	<LOQ (0.00750 ug/g) <LOQ (0.00750 ug/g) <LOQ (0.00750 ug/g) <LOQ (0.00750 ug/g)	<LOQ	NA
Yolk	Bird031-Gull Bird032-Gull Bird033-Gull	0.197 <LOQ (0.180 ug/g) 0.196	0.196 - 1 Outlier	0.528 0.00104	<LOQ (0.00750 ug/g) <LOQ (0.00750 ug/g) <LOQ (0.00750 ug/g)	<LOQ	NA

No PFOS qualitative confirmation performed. Identifications are preliminary.

LOQ = Limit of Quantitation

RSD = Relative Standard Deviation

RPD = Relative Percent Difference

Date Entered/Analyst: 10/23/99 LAC, 12/1/99 GML, 12/30/99 MMH

Date Verified/Analyst: 2/17/00 MMH

NA = Not Applicable

PFOS = Perfluorooctanesulfonate

PFHS = Perfluorohexanesulfonate

POAA = Perfluorooctanoate

PFOSA = Perfluorooctane sulfonamide

E

Study:
 Product Number(Test Substance):
 Matrix:
 Method/Revision:
 Analytical Equipment System Number:
 Instrument Software/Version:
 Filename:
 R-Squared Value:
 Slope:
 Y-Intercept:
 Dates of Extraction/Analyst:
 Dates of Analysis/Analyst:
 Date of Data Reduction/Analyst:

GEN030 MSU Environmental Samples
 Various
 Various Blood
 ETS-8-4.1 & ETS-8-5.1 using unextracted curves
 Soup020199
 Masslynx 3.3
 See Attachments
 See Attachments
 See Attachments
 See Attachments
 12/14/99 SAL/SRP/KK
 01/06/00, 01/07/00 MMH/IAS
 01/07/00, 01/10/00 IAS/MMH

Sample Data BLOOD

Group Desc	Sample #	Concentration of PFOS ug/mL or % Rec.	Mean PFOS ug/mL	RSD Std. Dev. MS/MSD RPD	Concentration of PFHS ug/mL or % Rec.	Mean PFHS ug/mL	RSD Std. Dev. MS/MSD RPD
Method Blk	MSU12129-H2Obk unfiltered 5-3 MSU12129-H2Obk filtered 5-4	<LOQ (0.00290 ug/mL) <LOQ (0.00290 ug/mL)	<LOQ	NA NA	<LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL)	<LOQ	NA NA
Matrix Blk	HMB12129-blood blk-5-1 HMB12129-blood blk-5-2 HMB12129-blood blk-5-3 HMB12129-blood blk-5-4 HMB12129-blood blk-5-5 HMB12129-blood blk-5-6* HMB12129-blood blk-5-7 HMB12129-blood blk-5-8* HMB12129-blood blk-5-9	0.0233 0.0248 0.0262 0.0243 <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL)			<LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL)		
QC	PBB-6255-250MSD-5-1-2 FSB-6255-250MSD-5-1-2 FSB-S009-250ppb MS-5-1-2 SSB-S009-250ppb MS-5-1-2 SSB-S5149-250ppb MS-5-1-2 HMB-FE52189-250 MS-5-1-2 HMB-FE52189-250 MSD-5-1-2	74% 91% -6% -5% 77% 65% -1% 77%			76% 77% -1% -1% 83% 77% 1% 74%		
Blood Northern Fur Seal Pups	P205 P206 P207 P208 P209 P210 P211 P212 P215 P217 P219 P220 P221 P222 P223 P224 P226 P229 P230	<LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL)			<LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL)		
Blood Northern Fur Seal Adult Females	M104 M105 M106 M107 M112 M115 M116 M118 M119 M122	<LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL)			<LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL)		
Blood Northern Fur Seal Subadult Males	S001 S002 S003 S006 S007 S008 S009	<LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL)			<LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL)		
Blood Northern Fur Seal	P234B P236A 368C P406 F411 98 CUKB 3* 98 CUKB 7 98 CUKB 9	<LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL) <LOQ (0.00579 ug/mL)			<LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL) <LOQ (0.00114 ug/mL)		

* Surrogate >50% deviation, not confirmed

I = May need to run all samples for PFOSA, interferent present in both analyses.

** PFOS NOT confirmed; MS transitions variation > 36%

Date Entered/By: 01/21/00 LAC

NE = Not Extracted

B = Lost during extraction

NA = Not Applicable

LOQ = Limit of Quantitation

PFOS = Perfluorooctanesulfonate

PFOSA = Perfluorooctane sulfonamide

PFHS = Perfluorohexanesulfonate

POAA = Perfluorooctanoate

Study:
 Product Number(Test Substance):
 Matrix:
 Method/Revision:
 Analytical Equipment System Number:
 Instrument Software/Version:
 Filename:
 R-Squared Value:
 Slope:
 Y-Intercept:
 Dates of Extraction/Analyst:
 Dates of Analysis/Analyst:
 Date of Data Reduction/Analyst:

GEN030 MSU Environmental Samples
 Various
 Various Blood
 ETS-8-4.1 & ETS-8-5.1 using unextracted curves
 Soup020199
 Masslynx 3.3
 See Attachments
 See Attachments
 See Attachments
 See Attachments
 12/14/99 SAL/SRP/KK
 01/06/00, 01/07/00 MMH/IAS
 01/07/00, 01/10/00 IAS/MMH

Sample Data

BLOOD

Group Data	Sample #	Concentration of POAA ug/mL or % Rec.	Mean POAA ug/mL	RSD Std. Dev. MS/MSD RPD	Concentration of PFOSA ug/mL or % Rec.	Mean PFOSA ug/mL	RSD Std. Dev. MS/MSD RPD
Method Blk	MSU12129-H2Obik unfiltered 5-3	<LOQ (0.00240 ug/mL)		NA	<LOQ (0.000625 ug/mL)		NA
	MSU12129-H2Obik filtered 5-4	<LOQ (0.00240 ug/mL)	<LOQ	NA	<LOQ (0.000625 ug/mL)	<LOQ	NA
Matrix Blk	HMB12129-blood blk-5-1	<LOQ (0.00240 ug/mL)			0.00193		
	HMB12129-blood blk-5-2	<LOQ (0.00240 ug/mL)			0.00190		
	HMB12129-blood blk-5-3	<LOQ (0.00240 ug/mL)			0.00225		
	HMB12129-blood blk-5-4	<LOQ (0.00240 ug/mL)			0.00189		
	HMB12129-blood blk-5-5	<LOQ (0.00958 ug/mL)			I		
	HMB12129-blood blk-5-6*	<LOQ (0.00958 ug/mL)			I		
	HMB12129-blood blk-5-7	<LOQ (0.00958 ug/mL)			I		
	HMB12129-blood blk-5-8	<LOQ (0.00958 ug/mL)		NA	I		NA
	HMB12129-blood blk-5-9	<LOQ (0.00958 ug/mL)	<LOQ	NA	I	I	NA
QC	PBB-6255-250MS-5-1-2	76%			50%		
	PBB-6255-250MSD-5-1-2	75%	76%	1%	47%	49%	6%
	FSB-S009-250ppb MS-5-1-2	0%			I		
	FSB-S009-250ppb MSD-5-1-2	1%	1%	36%	I	I	NA
	SSB-SSL49-250ppb MS-5-1-2	76%			I		
	SSB-SSL49-250ppb MSD-5-1-2	71%	74%	7%	I	I	NA
	HMB-FB52189-250 MS-5-1-2	1%			I		
	HMB-FB52189-250 MSD-5-1-2	70%	15%	195%	I	I	NA
Blood Northern Fur Seal Pups	P205	<LOQ (0.00958 ug/mL)			I		
	P206	<LOQ (0.00958 ug/mL)			I		
	P207	<LOQ (0.00958 ug/mL)			I		
	P208	<LOQ (0.00958 ug/mL)			I		
	P209	<LOQ (0.00958 ug/mL)			I		
	P210	<LOQ (0.00958 ug/mL)			I		
	P211	<LOQ (0.00958 ug/mL)			I		
	P212	<LOQ (0.00958 ug/mL)			I		
	P215	<LOQ (0.00958 ug/mL)			I		
	P217	<LOQ (0.00958 ug/mL)			I		
	P219	<LOQ (0.00958 ug/mL)			I		
	P220	<LOQ (0.00958 ug/mL)			I		
	P221	<LOQ (0.00958 ug/mL)			I		
	P222	<LOQ (0.00958 ug/mL)			I		
	P223	<LOQ (0.00958 ug/mL)			I		
	P224	<LOQ (0.00958 ug/mL)			I		
	P226	<LOQ (0.00958 ug/mL)			I		
	P229	<LOQ (0.00958 ug/mL)		NA	I		NA
	P230	<LOQ (0.00958 ug/mL)	<LOQ	NA	I	I	NA
Blood Northern Fur Seal Adult Females	M104	<LOQ (0.00958 ug/mL)			I		
	M105	<LOQ (0.00958 ug/mL)			I		
	M106	<LOQ (0.00958 ug/mL)			I		
	M107	<LOQ (0.00958 ug/mL)			I		
	M112	<LOQ (0.00958 ug/mL)			I		
	M115	<LOQ (0.00958 ug/mL)			I		
	M116	<LOQ (0.00958 ug/mL)			I		
	M118	<LOQ (0.00958 ug/mL)			I		
	M119	<LOQ (0.00958 ug/mL)		NA	I		NA
	M122	<LOQ (0.00958 ug/mL)	<LOQ	NA	I	I	NA
Blood Northern Fur Seal Subadult Males	S001	<LOQ (0.00958 ug/mL)			I		
	S002	<LOQ (0.00958 ug/mL)			I		
	S003	<LOQ (0.00958 ug/mL)			I		
	S006	<LOQ (0.00958 ug/mL)			I		
	S007	<LOQ (0.00958 ug/mL)			I		
	S008	<LOQ (0.00958 ug/mL)		NA	I		NA
	S009	<LOQ (0.00958 ug/mL)	<LOQ	NA	I	I	NA
Blood Northern Fur Seal	P298B	<LOQ (0.00958 ug/mL)			I		
	P236A	<LOQ (0.00958 ug/mL)			I		
	368C	<LOQ (0.00958 ug/mL)			I		
	P406	<LOQ (0.00958 ug/mL)			I		
	P411	<LOQ (0.00958 ug/mL)			I		
	98 CUKB 3*	<LOQ (0.00958 ug/mL)			I		
	98 CUKB 7	<LOQ (0.00958 ug/mL)		NA	I		NA
NA2 EDTA	98 CUKB 9	<LOQ (0.00958 ug/mL)	<LOQ	NA	I	I	NA

* Surrogate >50% deviation, not confirmed

I = May need to rerun all samples for PFOSA, interferent present in both analyses.

** PFOS NOT confirmed; MS transition variation > 30%

Date Entered/By: 01/21/00 LAC

NE = Not Extracted

E = Lost during extraction

NA = Not Applicable

LOQ = Limit of Quantitation

PFOS = Perfluorooctanesulfonate

PFOSA = Perfluorooctanesulfonamide

PFHS = Perfluorohexanesulfonate

POAA = Perfluorooctanoate

Study: GEN030 MSU Environmental Samples
 Product Number (Test Substance): Various
 Matrix: Various Blood
 Method/Revision: BTS-8-4.1 & ETS-8-5.1 using unextracted curves
 Analytical Equipment System Number: Soup020199
 Instrument Software/Version: Masslynx 3.3
 Filename: See Attachments
 R-Squared Value: See Attachments
 Slope: See Attachments
 Y-Intercept: See Attachments
 Dates of Extraction/Analyst: 12/14/99 SAL/SRP/KKK
 Dates of Analysis/Analyst: 01/06/00, 01/07/00 MMH/IAS
 Date of Data Reduction/Analyst: 01/07/00, 01/10/00 IAS/MMH

Sample Data

BLOOD

Group Desc	Sample #	PFOS Verified	Concentration of PFOS ug/mL or % Rec.	Mean PFOS ug/mL	RSD Std. Dev. MS/MSD RPD	Concentration of PFHS ug/mL or % Rec.	Mean PFHS ug/mL	RSD Std. Dev. MS/MSD RPD
Method Blk	MSU12129-H2Obk unfiltered 5-3	NA	<LOQ (0.00290 ug/mL)	<LOQ	NA	<LOQ (0.00114 ug/mL)	<LOQ	NA
	MSU12129-H2Obk filtered 5-4	NA	<LOQ (0.00290 ug/mL)	<LOQ	NA	<LOQ (0.00114 ug/mL)	<LOQ	NA
Matrix Blk	HMB12129-blood blk-5-1	NA	0.0253			<LOQ (0.00114 ug/mL)		
	HMB12129-blood blk-5-2	NA	0.0248			<LOQ (0.00114 ug/mL)		
	HMB12129-blood blk-5-3	NA	0.0262			<LOQ (0.00114 ug/mL)		
	HMB12129-blood blk-5-4	NA	0.0243			<LOQ (0.00114 ug/mL)		
	HMB12129-blood blk-5-5	NA	<LOQ (0.0579 ug/mL)			<LOQ (0.0114 ug/mL)		
	HMB12129-blood blk-5-6*	NA	<LOQ (0.0579 ug/mL)			<LOQ (0.0114 ug/mL)		
	HMB12129-blood blk-5-7	NA	<LOQ (0.0579 ug/mL)			<LOQ (0.0114 ug/mL)		
	HMB12129-blood blk-5-8	NA	<LOQ (0.0579 ug/mL)			<LOQ (0.0114 ug/mL)		
	HMB12129-blood blk-5-9	NA	<LOQ (0.0579 ug/mL)	<LOQ - 4 outliers	NA 0.000790	<LOQ (0.0114 ug/mL)	<LOQ	NA NA
QC	PBB-6255-250MS-5-1-2	NA	74%			76%		
	PBB-6255-250MSD-5-1-2	NA	91%	82%	21%	77%	76%	2%
	FSB-S009-250ppb MS-5-1-2	NA	-6%	-6%		-1%		
	FSB-S009-250ppb MSD-5-1-2	NA	-5%	-6%	25%	-1%	-1%	20%
	SSB-SSL49-250ppb MS-5-1-2	NA	77%			83%		
	SSB-SSL49-250ppb MSD-5-1-2	NA	65%	71%	17%	77%	80%	9%
	HMB-FE52189-250 MS-5-1-2	NA	-1%			1%		
	HMB-FE52189-250 MSD-5-1-2	NA	77%	38%	208%	74%	37%	196%
Blood Polar Bear	6255 (Heparin)	X	0.0518			<LOQ (0.00114 ug/mL)		
	20436 (NA2 EDTA)	X	0.0381			<LOQ (0.00114 ug/mL)		
	20467 (NA2 EDTA)	X	0.0358			<LOQ (0.00114 ug/mL)		
	20468 (EDTA)	X	0.0315			<LOQ (0.00114 ug/mL)		
	20470	X	0.0281			<LOQ (0.00114 ug/mL)		
	20472	X	0.0327			<LOQ (0.00114 ug/mL)		
	20473*	X	0.0309			<LOQ (0.00114 ug/mL)		
	20474 (NA2 EDTA)	X	0.0256			<LOQ (0.00114 ug/mL)		
	20475 (NA2 EDTA)	X	<LOQ (0.00290 ug/mL)			<LOQ (0.00114 ug/mL)		
	20476 (NA2 EDTA)	X	0.0345			<LOQ (0.00114 ug/mL)		
	20477	X	0.0272			<LOQ (0.00114 ug/mL)		
	20485 (Heparin)	X	0.0429			<LOQ (0.00114 ug/mL)		
	20486 (Heparin)	X	0.0316		0.214	<LOQ (0.00114 ug/mL)		NA
	20487 (Heparin)	X	<LOQ (0.00290 ug/mL)	0.0342 - 2 outliers	0.00731	<LOQ (0.00114 ug/mL)	<LOQ	NA
Blood Stellar Sealion	SSL49 (7.2 mg K2 EDTA)	NA	<LOQ (0.00579 ug/mL)			<LOQ (0.0114 ug/mL)		
	SSL50 (7.2 mg K2 EDTA)	NA	<LOQ (0.00579 ug/mL)			<LOQ (0.0114 ug/mL)		
	SSL51 (7.2 mg K2 EDTA)	NA	<LOQ (0.00579 ug/mL)			<LOQ (0.0114 ug/mL)		
	SSL52 (7.2 mg K2 EDTA)	NA	<LOQ (0.00579 ug/mL)			<LOQ (0.0114 ug/mL)		
	SSL53 (7.2 mg K2 EDTA)	NA	<LOQ (0.00579 ug/mL)			<LOQ (0.0114 ug/mL)		
	SSL54 (7.2 mg K2 EDTA)	NA	<LOQ (0.00579 ug/mL)			<LOQ (0.0114 ug/mL)		
	SSL55 (5.4 mg K2 EDTA)	NA	<LOQ (0.00579 ug/mL)			<LOQ (0.0114 ug/mL)		
	SSL56 (5.4 mg K2 EDTA)*	NA	<LOQ (0.00579 ug/mL)			<LOQ (0.0114 ug/mL)		
	SSL57 (5.4 mg K2 EDTA)	NA	<LOQ (0.00579 ug/mL)			<LOQ (0.0114 ug/mL)		
	SSL58 (5.4 mg K2 EDTA)*	NA	<LOQ (0.00579 ug/mL)			<LOQ (0.0114 ug/mL)		
	SSL59 (5.4 mg K2 EDTA)	NA	<LOQ (0.00579 ug/mL)		NA	<LOQ (0.0114 ug/mL)		NA
	SSL60 (5.4 mg K2 EDTA)*	NA	<LOQ (0.00579 ug/mL)	<LOQ	NA	<LOQ (0.0114 ug/mL)	<LOQ	NA

* = May need to rerun all samples for PFOSA, interferent present in both analyses.

** Surrogate >50% deviation confirmed

** Surrogate >50% deviation, not confirmed

** PFOS NOT confirmed; MS transitions variation > 30%

Date Entered/By: 1/13/00, 01/21/00 MMH/LAC

Date Verified/By: 02/10/00 kjh

NE = Not Extracted

E = Lost during extraction

NA = Not Applicable

NV = Not Verified

LOQ = Limit of quantitation

PFOS = Perfluorooctanesulfonate

PFOSA = Perfluorooctanesulfonamide

PFHS = Perfluorohexanesulfonate

POAA = Perfluorooctanoate

Study:
Product Number(Test Substance):
Matrix:
Method/Revision:
Analytical Equipment System Number:
Instrument Software/Version:
Filename:
R-Squared Value:
Slope:
Y-Intercept:
Dates of Extraction/Analyst:
Dates of Analysis/Analyst:
Date of Data Reduction/Analyst:

GEN030 MSU Environmental Samples
Various
Various Blood
ETS-B-4.1 & ETS-B-5.1 using unextracted-curves
Soup020199
Masslynx 3.3
See Attachments
See Attachments
See Attachments
See Attachments
12/14/99 SAL/SRP/KCK
01/06/00, 01/07/00 MMH/LAS
01/07/00, 01/10/00 IAS/MMH

Sample Data

BLOOD

Group Dose	Sample #	Concentration of POAA ug/mL or % Rec.	Mean POAA ug/mL	RSD Std. Dev. MS/MSD RPD	Concentration of PFOSA ug/mL or % Rec.	Mean PFOSA ug/mL	RSD Std. Dev. MS/MSD RPD
Method Blk	MSU12129-H2Oblk unfiltered 5-3 MSU12129-H2Oblk filtered 5-4	<LOQ (0.00240 ug/mL) <LOQ (0.00240 ug/mL)	<LOQ	NA NA	<LOQ (0.000625 ug/mL) <LOQ (0.000625 ug/mL)	<LOQ	NA NA
Matrix Blk	HMB12129-blood blk-5-1 HMB12129-blood blk-5-2 HMB12129-blood blk-5-3 HMB12129-blood blk-5-4 HMB12129-blood blk-5-5 HMB12129-blood blk-5-6* HMB12129-blood blk-5-7 HMB12129-blood blk-5-8 HMB12129-blood blk-5-9	<LOQ (0.00240 ug/mL) <LOQ (0.00240 ug/mL) <LOQ (0.00240 ug/mL) <LOQ (0.00240 ug/mL) <LOQ (0.00958 ug/mL) <LOQ (0.00958 ug/mL) <LOQ (0.00958 ug/mL) <LOQ (0.00958 ug/mL) <LOQ (0.00958 ug/mL)	<LOQ	NA NA	<LOQ (0.000625 ug/mL) <LOQ (0.000625 ug/mL) <LOQ (0.000625 ug/mL) <LOQ (0.000625 ug/mL) I I I I I	I	NA NA
QC	PBB-6255-250MSD-5-1-2 PBB-6255-250MSD-5-1-2 FSB-S009-250ppb MS-5-1-2 FSB-S009-250ppb MSD-5-1-2 SSB-SSL49-250ppb MS-5-1-2 SSB-SSL49-250ppb MSD-5-1-2 HMB-FE52189-250 MS-5-1-2 HMB-FE52189-250 MSD-5-1-2	76% 75% 0% 1% 76% 71% 1% 70%	76% 76% 1% 1% 74% 74% 35%	1% 1% 36% 7% 195%	50% 47% I I I I I I	49% I I I I	6% NA NA NA NA
Blood Polar Bear	6255 (Heparin) 20436 (NA2 EDTA) 20467 (NA2 EDTA) 20468 (EDTA) 20470 20472 20473* 20474 (NA2 EDTA) 20475 (NA2 EDTA) 20476 (NA2 EDTA) 20477 20485 (Heparin) 20486 (Heparin) 20487 (Heparin)	<LOQ (0.00240 ug/mL) <LOQ (0.00240 ug/mL) <LOQ (0.00240 ug/mL) <LOQ (0.00240 ug/mL) <LOQ (0.00240 ug/mL) <LOQ (0.00240 ug/mL) <LOQ (0.00240 ug/mL) <LOQ (0.00240 ug/mL) <LOQ (0.00240 ug/mL) <LOQ (0.00240 ug/mL) <LOQ (0.00240 ug/mL) <LOQ (0.00240 ug/mL) <LOQ (0.00240 ug/mL) <LOQ (0.00240 ug/mL)	<LOQ	NA NA	<LOQ (0.000625 ug/mL) <LOQ (0.000625 ug/mL) <LOQ (0.000625 ug/mL) <LOQ (0.000625 ug/mL) <LOQ (0.000625 ug/mL) <LOQ (0.000625 ug/mL) <LOQ (0.000625 ug/mL) <LOQ (0.000625 ug/mL) <LOQ (0.000625 ug/mL) <LOQ (0.000625 ug/mL) <LOQ (0.000625 ug/mL) <LOQ (0.000625 ug/mL) <LOQ (0.000625 ug/mL) <LOQ (0.000625 ug/mL)	<LOQ	NA NA
Blood Stellar Sealion	SSL49 (7.2 mg K2 EDTA) SSL50 (7.2 mg K2 EDTA) SSL51 (7.2 mg K2 EDTA) SSL52 (7.2 mg K2 EDTA) SSL53 (7.2 mg K2 EDTA) SSL54 (7.2 mg K2 EDTA) SSL55 (5.4 mg K2 EDTA) SSL56 (5.4 mg K2 EDTA)** SSL57 (5.4 mg K2 EDTA) SSL58 (5.4 mg K2 EDTA)* SSL59 (5.4 mg K2 EDTA) SSL60 (5.4 mg K2 EDTA)*	<LOQ (0.00958 ug/mL) <LOQ (0.00958 ug/mL) <LOQ (0.00958 ug/mL) <LOQ (0.00958 ug/mL) <LOQ (0.00958 ug/mL) <LOQ (0.00958 ug/mL) <LOQ (0.00958 ug/mL) <LOQ (0.00958 ug/mL) <LOQ (0.00958 ug/mL) <LOQ (0.00958 ug/mL) <LOQ (0.00958 ug/mL) <LOQ (0.00958 ug/mL)	<LOQ	NA NA	I I I I I I I I I I I I	I	NA NA

I = May need to rerun all samples for PFOSA, interferent present in both analyses

** Surrogate >50% deviation confirmed

* Surrogate >30% deviation, not confirmed

** PFOS NOT confirmed; MS transitions variation > 30%

Date Entered/By: 1/13/00, 01/21/00 MMH/LAC

Date Verified/By: 02/10/00 kjh

NE = Not Extracted

E = Lost during extraction

NA = Not Applicable

NV = Not Verified

LOQ = Limit of quantitation

PFOS = Perfluorooctanesulfonate

PFOSA = Perfluorooctanesulfonamide

PFHS = Perfluorohexanesulfonate

POAA = Perfluorooctanone



Wilson Jones - Quick Reference Index System

© 1991 Wilson Jones Company

3M_MN01666030

2814.0034

FACT-GEN-030
MSU Environmental Samples

Study:	GEN030 MSU Environmental Samples
Product Number(Test Substance):	Various
Matrix:	Various
Method/Revision:	ETS-8-6.0 & ETS-8-7.0
Analytical Equipment System Number:	Soup 020199, Amelia 062498
Instrument Software/Version:	Masslynx 3.3
Date of Extraction/Analyst:	12/12/99 SAL/KK/SRP/CSH
Date of Analysis/Analyst:	01/10/00 MMH
Date of Data Reduction/Analyst:	01/12/00 LAS

Sample Data

MUSCLE

Group Dose	Sample #	PFOS Verified	Concentration of PFOS ug/g or % Rec.	Mean PFOS ug/g	RSD Std. Dev. MS/MSD RPD	Concentration of PFHS ug/g or % Rec.	Mean PFHS ug/g	RSD Std. Dev. MS/MSD RPD
Method Blk	MSU12149-H2Ounfil 5-1*	NA	0.0163		NA	<LOQ (0.0342)		NA
Unfiltered water	MSU12149-H2Ounfil 5-2*	NA	<LOQ (0.00696)	NA	NA	<LOQ (0.0342)	<LOQ	NA
Method Blk	MSU12149-H2Ofil 5-1*	NA	<LOQ (0.00696)		NA	<LOQ (0.0342)		NA
Filtered water	MSU12149-H2Ofil 5-2*	NA	0.0257	NA	NA	<LOQ (0.0342)	<LOQ	NA
Matrix Blk	MSU12149-Fishblk 5-1	NA	0.00948			<LOQ (0.0342)		NA
Fish Liver	MSU12149-Fishblk 5-2	NA	0.00985		37.2	<LOQ (0.0342)		NA
	MSU12149-Fishblk 5-3*	NA	0.00454	0.00796	0.00296	<LOQ (0.0342)	<LOQ	NA
QC 250 ug/g	CPM-B1N7-MS 5-1-2	NA	59%			60%		
	CPM-B2N6-MS 5-1-2	NA	31%	45%	63%	45%	53%	28%
	CSM-1999030-03-01-MS 5-1-2	NA	160%			148%		
	CSM-1999030-03-01-MSD 5-1-2	NA	127%	143%	23%	183%	166%	21%
	LWM-1999029-16-MS 5-1-2	NA	67%			129%		
	LWM-1999029-11-MS 5-1-3	NA	122%	95%	59%	130%	130%	0%
	BTM-1999040-06-MS 5-1-2	NA	148%			144%		
	BTM-1999040-10-MS 5-1-2	NA	166%	157%	11%	168%	156%	15%
Muscle Carp	B1N1	X	0.101			<LOQ (0.0342)		
	B1N2	X	0.0784			<LOQ (0.0342)		
	B1N7	X	0.0905			<LOQ (0.0342)		
	B1N10	X	0.0878			<LOQ (0.0342)		
	B2N2	X	0.105			<LOQ (0.0342)		
	B2N6	X	0.0894			<LOQ (0.0342)		
	B2N8	X	0.0596			<LOQ (0.0342)		
	B2N10	X	0.0838			<LOQ (0.0342)		
	Carp1*	X	0.297		64.2	<LOQ (0.0342)		NA
	Carp2*	X	0.243	0.124	0.0793	<LOQ (0.0342)	<LOQ	NA
Muscle Chinook Salmon	1999030-01	NA	0.189			<LOQ (0.0342)		
	1999030-02	NA	0.126			<LOQ (0.0342)		
	1999030-02-01	X	<LOQ (0.00696)			<LOQ (0.0342)		
	1999030-02-04	X	0.113			<LOQ (0.0342)		
	1999030-03-01	X	0.0514		0.524	<LOQ (0.0342)		NA
	1999030-03-04	X	0.0573	0.107 - 1 outlier	0.0562	<LOQ (0.0342)	<LOQ	NA
Muscle Lake Whitefish	1999029-11*	X	0.168			<LOQ (0.0342)		
	1999029-12*	X	0.130			<LOQ (0.0342)		
	1999029-13	X	0.0967			<LOQ (0.0342)		
	1999029-14	X	0.0983		26.4	<LOQ (0.0342)		NA
	1999029-16	X	0.1639	0.132	0.0348	<LOQ (0.0342)	<LOQ	NA
Muscle Brown Trout	1999040-01	X	<LOQ (0.00696)			<LOQ (0.0342)		
	1999040-02 **	X **	<LOQ (0.00696)			<LOQ (0.0342)		
	1999040-03	NA	<LOQ (0.00696)			<LOQ (0.0342)		
	1999040-04	NA	<LOQ (0.00696)			<LOQ (0.0342)		
	1999040-05	X	<LOQ (0.00696)			<LOQ (0.0342)		
	1999040-06	X	<LOQ (0.00696)			<LOQ (0.0342)		
	1999040-07	NA	<LOQ (0.00696)			<LOQ (0.0342)		
	1999040-08	X	0.0460			<LOQ (0.0342)		
	1999040-09	NA	<LOQ (0.00696)		NA	<LOQ (0.0342)		NA
	1999040-10	NA	<LOQ (0.00696)	<LOQ - 1 outlier	NA	<LOQ (0.0342)	<LOQ	NA

* High (>50%) surrogate deviations

** PFOS NOT confirmed; MS transitions variation > 30%

Date Entered/Analyst: 01/24/00, 01/25/00, 01/28/00 LAC

Date Verified/Analyst: 02/10/00 kjh

NE = Not Extracted

E = Lost during extraction

NA = Not Applicable

LOQ = Limit of Quantitation

X = Verified PFOS concentration

PFOS = Perfluorooctanesulfonate

PFOSA = Perfluorooctane sulfonamide

PFHS = Perfluorohexanesulfonate

POAA = Perfluorooctanamide

FACT-GEN-030
MSU Environmental Samples

Study:
Product Number(Test Substance):
Matrix:
Method/Revision:
Analytical Equipment System Number
Instrument Software/Version
Date of Extraction/Analyst:
Date of Analysis/Analyst:
Date of Data Reduction/Analyst:
Sample Data

GEN030 MSU Environmental Samples
Various
Various
ETS-8-6.0 & ETS-8-7.0
Soup 020199, Amelia 062498
Masslynx 3.3
12/12/99 SAL/KJ/SRP/CSH
01/10/00 MMH
01/12/00 IAS

MUSCLE

Group Dose	Sample #	Concentration of POAA ug/g or % Rec.	Mean POAA ug/g	RSD Std. Dev. MS/MSD RPD	Concentration of PFOSA ug/g or % Rec.	Mean PFOSA ug/g	RSD Std. Dev. MS/MSD RPD
Method Blk	MSU12149-H2Oumfil 5-1*	<LOQ (0.0359)		NA	<LOQ (0.0188)		NA
Unfiltered water	MSU12149-H2Oumfil 5-2*	<LOQ (0.0359)	<LOQ	NA	<LOQ (0.0188)	<LOQ	NA
Method Blk	MSU12149-H2Oumfil 5-1*	<LOQ (0.0359)		NA	<LOQ (0.0188)		NA
Filtered water	MSU12149-H2Oumfil 5-2*	<LOQ (0.0359)	<LOQ	NA	<LOQ (0.0188)	<LOQ	NA
Matrix Blk	MSU12149-Fishblk 5-1	<LOQ (0.0359)		NA	<LOQ (0.0188)		NA
Fish Liver	MSU12149-Fishblk 5-2	<LOQ (0.0359)		NA	<LOQ (0.0188)		NA
	MSU12149-Fishblk 5-3*	<LOQ (0.0359)	<LOQ	NA	<LOQ (0.0188)	<LOQ	NA
QC 250 ng/g	CPM-B1N7-MS 5-1-2	6%			11%		
	CPM-B2N6-MS 5-1-2	39%	23%	148%	13%	12%	19%
	CSM-1999030-03-01-MS 5-1-2	139%			77%		
	CSM-1999030-03-01-MSD 5-1-2	178%	158%	25%	69%	73%	11%
	LWM-1999029-16-MS 5-1-2	146%			61%		
	LWM-1999029-11-MS 5-1-3	105%	125%	33%	76%	68%	22%
	BTM-1999040-09-MS 5-1-2	147%			80%		
	BTM-1999040-10-MS 5-1-2	152%	150%	3%	87%	83%	8%
Muscle Carp	B1N1	<LOQ (0.0359)			<LOQ (0.0188)		
	B1N2	<LOQ (0.0359)			<LOQ (0.0188)		
	B1N7	<LOQ (0.0359)			<LOQ (0.0188)		
	B1N10	<LOQ (0.0359)			<LOQ (0.0188)		
	B2N2	<LOQ (0.0359)			<LOQ (0.0188)		
	B2N6	<LOQ (0.0359)			<LOQ (0.0188)		
	B2N8	<LOQ (0.0359)			<LOQ (0.0188)		
	B2N10	<LOQ (0.0359)			<LOQ (0.0188)		
	Carp1*	<LOQ (0.0359)		NA	<LOQ (0.0188)		NA
Muscle Chinook Salmon	Carp2*	<LOQ (0.0359)	<LOQ	NA	<LOQ (0.0188)	<LOQ	NA
	1999030-01	<LOQ (0.0359)			<LOQ (0.0188)		
	1999030-02	<LOQ (0.0359)			<LOQ (0.0188)		
	1999030-02-01	<LOQ (0.0359)			<LOQ (0.0188)		
	1999030-02-04	<LOQ (0.0359)			<LOQ (0.0188)		
	1999030-03-01	<LOQ (0.0359)		NA	<LOQ (0.0188)		NA
Muscle Lake Whitefish	1999030-03-04	<LOQ (0.0359)	<LOQ	NA	<LOQ (0.0188)	<LOQ	NA
	1999029-11*	<LOQ (0.0359)			<LOQ (0.0188)		
	1999029-12*	<LOQ (0.0359)			<LOQ (0.0188)		
	1999029-13	<LOQ (0.0359)			<LOQ (0.0188)		
	1999029-14	<LOQ (0.0359)		NA	<LOQ (0.0188)		NA
	1999029-16	<LOQ (0.0359)	<LOQ	NA	<LOQ (0.0188)	<LOQ	NA
Muscle Brown Trout	1999040-01	<LOQ (0.0359)			<LOQ (0.0188)		
	1999040-02 **	<LOQ (0.0359)			<LOQ (0.0188)		
	1999040-03	<LOQ (0.0359)			<LOQ (0.0188)		
	1999040-04	<LOQ (0.0359)			<LOQ (0.0188)		
	1999040-05	<LOQ (0.0359)			<LOQ (0.0188)		
	1999040-06	<LOQ (0.0359)			<LOQ (0.0188)		
	1999040-07	<LOQ (0.0359)			<LOQ (0.0188)		
	1999040-08	<LOQ (0.0359)			<LOQ (0.0188)		
	1999040-09	<LOQ (0.0359)		NA	<LOQ (0.0188)		NA
	1999040-10	<LOQ (0.0359)	<LOQ	NA	<LOQ (0.0188)	<LOQ	NA

* High (>50%) surrogate deviations

** PFOS NOT confirmed; MS transitions variation > 30%

Date Entered/Analyst: 01/24/00, 01/25/00, 01/28/00 LAC

Date Verified/Analyst: 02/10/00 kjh

NE = Not Extracted

E = Lost during extraction

NA = Not Applicable

LOQ = Limit of Quantitation

X = Verified PFOS concentration

PFOS = Perfluorooctanesulfonate

PFOSA = Perfluorooctane sulfonamide

PFHS = Perfluorohexanesulfonate

POAA = Perfluorooctanoate

FACT-GEN-030
MSU Environmental Samples

Study:
Product Number/Test Substance):
Matrix:
Method/Revision:
Analytical Equipment System Number:
Instrument Software/Version:
Date of Extraction/Analyst:
Date of Analysis/Analyst:
Date of Data Reduction/Analyst:
Sample Data

GEN030 MSU Environmental Samples
Various
Various
BTS-8-6.0 & BTS-8-7.0
Amelia 062498
Mamlynx 3.3
12/12/99 SAL/KK/SRP/CSH
01/10/00 MMH
01/11/00 IAS

MUSCLE and EGGS

Group Dose	Sample #	PFOS Verified	Concentration of PFOS ug/g or % Rec.	Mean PFOS ug/g	RSD Std. Dev. MS/MSD RPD	Concentration of PFOS ug/g or % Rec.	Mean PFOS ug/g	RSD Std. Dev. MS/MSD RPD
Method Blk	MSU12159-H2OBlk-unfiltered-5-9	NA	<LOQ (0.0174)	<LOQ	NA	<LOQ (0.0342)	<LOQ	NA
Water	MSU12159-H2OBlk-filtered-5-9	NA	<LOQ (0.0174)	<LOQ	NA	<LOQ (0.0342)	<LOQ	NA
Matrix Blk	EggBlk-5-1	NA	<LOQ (0.0174)	<LOQ	NA	<LOQ (0.0342)	<LOQ	NA
Egg	EggBlk-5-2	NA	<LOQ (0.0174)	<LOQ	NA	<LOQ (0.0342)	<LOQ	NA
	EggBlk-5-3	NA	<LOQ (0.0174)	<LOQ	NA	<LOQ (0.0342)	<LOQ	NA
QC 250 ng/g	FGW-ALSWCR-TD2 6/25/98-MS-5-1-2	X	44%	46%	8%	16%	16%	3%
	FGW-ALSWCR-TD1 6/25/98-MSD-5-1-2	X	47%	46%	8%	16%	16%	3%
	FGW-AL118-HP26/25/98-MS-5-1-2	X	13%	14%	8%	10%	11%	13%
	FGW-AL118-HP26/25/98-MSD-5-1-2	X	15%	14%	8%	12%	11%	13%
	LWE-19999029-13-MS-5-1-2	X	146%	125%	33%	30%	33%	16%
	LWE-19999029-13-MSD-5-1-2	X	104%	125%	33%	35%	33%	16%
Carp	BTE-19999040-01-MS-5-1-2	X	121%	144%	32%	57%	57%	1%
	BTE-19999040-01-MSD-5-1-2	X	167%	144%	32%	57%	57%	1%
	Diet 1 (Carp5) **	X **	<LOQ (0.0174)	0.0272 - 2 outliers	NA	<LOQ (0.0342)	<LOQ	NA
	Diet 2 (Carp6)	X	<LOQ (0.0174)		NA	<LOQ (0.0342)	<LOQ	NA
Frog Muscle Wholebody	Diet 3 (Carp4)	X	0.0267		NA	<LOQ (0.0342)	<LOQ	NA
	Diet 4 (Carp3)	X	0.0278		NA	<LOQ (0.0342)	<LOQ	NA
	AL-118-V0Y 08/26/98 **	X **	0.00243			<LOQ (0.0342)		
	ALSWCR-TD 06/03/98 **	X **	<LOQ (0.0174)			<LOQ (0.0342)		
	ALSWCR-TD2 06/25/99 **	X **	<LOQ (0.0174)			<LOQ (0.0342)		
	KZCKDM-JUV 08/26/98 **	X **	<LOQ (0.0174)			<LOQ (0.0342)		
	KZCKDM-TD 06/05/98 **	X **	0.0216			<LOQ (0.0342)		
	KZCKDM-TD-2 06/25/98 **	X **	<LOQ (0.0174)			<LOQ (0.0342)		
Green Frog Eggs	SJ0002-TD-2 06/05/98 **	X **	<LOQ (0.0174)	<LOQ - 2 outliers	NA	<LOQ (0.0342)	<LOQ	NA
	SJ0002-TD-2 06/25/98 **	X **	<LOQ (0.0174)		NA	<LOQ (0.0342)	<LOQ	NA
	AL-118-HP29 06/25/98	X	<LOQ (0.0174)			<LOQ (0.0342)		
	AL-118-HP94 06/25/98	X	<LOQ (0.0174)			<LOQ (0.0342)		
Lake Whitefish Eggs	SJ0001 06/03/98 **	X **	<LOQ (0.0174)		NA	<LOQ (0.0342)		NA
	SJ0001 06/05/98	X	<LOQ (0.0174)	<LOQ	NA	<LOQ (0.0342)	<LOQ	NA
	1999029-13	X	0.145		63.5	<LOQ (0.0342)		NA
Brown Trout Eggs	1999029-14	X	0.381	0.263	0.167	<LOQ (0.0342)	<LOQ	NA
	1999040-01	X	0.0749			<LOQ (0.0342)		
	1999040-04	X	0.0675		21.1	<LOQ (0.0342)		NA
	1999040-06	X	0.0488	0.0637	0.0134	<LOQ (0.0342)	<LOQ	NA

* High (>50%) surrogate deviations

** PFOS NOT confirmed; MS transitions variation > 30%

Date Entered/Analyst: 01/21/00, 01/24/00 LAC

Date Verified/Analyst: 02/10/00 kjh

NE = Not Extracted

E = Lost during extraction

NA = Not Applicable

LOQ = Limit of Quantitation

X = Verified PFOS concentration

PFOS = Perfluorooctanesulfonate

PFOSA = Perfluorooctane sulfonamide

PFHS = Perfluorohexanesulfonate

FOAA = Perfluorooctanoate

Study:
Product Number/Test Substance:
Matrix:
Method/Revision:
Analytical Equipment System Number:
Instrument Software/Version:
Date of Extraction/Analyst:
Date of Analysis/Analyst:
Date of Data Reduction/Analyst:
Sample Data

GEN030 MSU Environmental Samples
Various
Various
ETS-8-6.0 & ETS-8-7.0
Amelia 062498
Masslynx 3.3
12/12/99 SAL/KK/SRP/CSH
01/10/00 MMH
01/11/00 IAS

MUSCLE and EGGS

Group Dose	Sample #	Concentration of POAA ug/g or % Rec.	Mean POAA ug/g	RSD Std. Dev. MS/MSD RPD	Concentration of PFOSA ug/g or % Rec.	Mean PFOSA ug/g	RSD Std. Dev. MS/MSD RPD
Method Blk	MSU12159-H2OBK-unfiltered-5-9	<LOQ (0.0359)	<LOQ	NA	<LOQ (0.0188)	<LOQ	NA
Water	MSU12159-H2OBK-filtered-5-9	<LOQ (0.0359)	<LOQ	NA	<LOQ (0.0188)	<LOQ	NA
Matrix Blk	EggBlk-5-1	<LOQ (0.0359)	<LOQ	NA	<LOQ (0.0188)	<LOQ	NA
Egg	EggBlk-5-2	<LOQ (0.0359)	<LOQ	NA	<LOQ (0.0188)	<LOQ	NA
	EggBlk-5-3	<LOQ (0.0359)	<LOQ	NA	<LOQ (0.0188)	<LOQ	NA
QC 250 ug/g	PGW-ALSWCR-TD1 6/25/98-MS-5-1-2	34%			49%		
	PGW-ALSWCR-TD1 6/25/98-MSD-5-1-2	33%	34%	4%	49%	49%	1%
	FGB-AL118-HP26/25/98-MS-5-1-2	22%			46%		
	FGB-AL118-HP26/25/98-MSD-5-1-2	26%	24%	15%	54%	50%	16%
	LWE-19999029-13-MS-5-1-2	90%			68%		
	LWE-19999029-13-MSD-5-1-2	83%	86%	9%	64%	66%	6%
Carp	BTE-19999040-01-MS-5-1-2	98%			104%		
	BTE-19999040-01-MSD-5-1-2	112%	105%	13%	108%	106%	4%
	Diet 1 (Carp5) **	<LOQ (0.0359)			<LOQ (0.0188)		
	Diet 2 (Carp6)	<LOQ (0.0359)			<LOQ (0.0188)		
Frog Muscle Wholebody	Diet 3 (Carp4)	<LOQ (0.0359)		NA	<LOQ (0.0188)		NA
	Diet 4 (Carp3)	<LOQ (0.0359)	<LOQ	NA	<LOQ (0.0188)	<LOQ	NA
	AL-118-YOY 06/26/98 **	<LOQ (0.0180)			<LOQ (0.0188)		
	ALSWCR-TD 06/03/98 **	<LOQ (0.0180)			<LOQ (0.0188)		
	ALSWCR-TD1 06/25/99 **	<LOQ (0.0180)			<LOQ (0.0188)		
	KZCKDM-JUV 06/26/98 **	<LOQ (0.0180)			<LOQ (0.0188)		
	KZCKDM-TD 06/05/98 **	<LOQ (0.0180)			<LOQ (0.0188)		
	KZCKDM-TD-2 06/25/98 **	<LOQ (0.0180)			<LOQ (0.0188)		
Green Frog Eggs	SJ0002-TD-2 06/05/98 **	<LOQ (0.0180)		NA	<LOQ (0.0188)		NA
	SJ0002-TD-2 06/25/98 **	<LOQ (0.0180)	<LOQ	NA	<LOQ (0.0188)	<LOQ	NA
	AL-118-HP89 06/25/98	<LOQ (0.0180)			<LOQ (0.0188)		
	AL-118-HP94 06/25/98	<LOQ (0.0180)			<LOQ (0.0188)		
Lake Whitefish Eggs	SJ0001 06/03/98 **	<LOQ (0.0180)		NA	<LOQ (0.0188)		NA
	SJ0001 06/05/98	<LOQ (0.0180)	<LOQ	NA	<LOQ (0.0188)	<LOQ	NA
	1999029-13	<LOQ (0.0180)		NA	<LOQ (0.0188)		NA
Brown Trout Eggs	1999029-14	<LOQ (0.0180)	<LOQ	NA	<LOQ (0.0188)	<LOQ	NA
	1999040-01	<LOQ (0.0180)			<LOQ (0.0188)		
	1999040-04	<LOQ (0.0180)		NA	<LOQ (0.0188)		NA
	1999040-06	<LOQ (0.0180)	<LOQ	NA	<LOQ (0.0188)	<LOQ	NA

* High (>50%) surrogate deviations

** PFOS NOT confirmed; MS transitions variation > 30%

Date Entered/Analyst: 01/21/00, 01/24/00 LAC

Date Verified/Analyst: 02/10/00 kjh

NE = Not Extracted

E = Lost during extraction

NA = Not Applicable

LOQ = Limit of Quantitation

X = Verified PFOS concentration

PFOS = Perfluorooctanesulfonate

PFOSA = Perfluorooctane sulfonamide

PFHS = Perfluorohexanesulfonate

POAA = Perfluorooctanoate

FACT-GEN-030
MSU Environmental Samples

Study	GEN030 MSU Environmental Samples
Product Number(Test Substance)	Various
Matrix	Various
Method/Revision	ETS-8-6.0 & ETS-8-7.0
Analytical Equipment System Number	Soup 020199, Amelia 062498
Instrument Software/Version	Masslynx 3.3
Date of Extraction/Analyst	12/12/99 SAL/KK/SRP/CSH
Date of Analysis/Analyst	12/17/99, 12/20/99, 12/28/99, 12/29/99, 01/03/00, 01/06/00 IAS/MMH
Date of Data Reduction/Analyst	12/20/99, 12/21/99, 12/22/99, 12/30/99, 01/03/00, 01/05/00, 01/07/00 MMH/IAS

Sample Data

Miscellaneous Liver

Group Dose	Sample #	PFOS Verified	Concentration of PFOS ug/g or % Rec.	Mean PFOS ug/g	RSD Std. Dev. MS/MSD RPD	Concentration of PFHS ug/g or % Rec.	Mean PFHS ug/g	RSD Std. Dev. MS/MSD RPD
Method Blk Unfiltered water	MSU12129-H2OBik-unfiltered-5-1	NA	NA			NA		
	MSU12129-H2OBik-unfiltered-5-2	NA	NA			NA		
	MSU12129-H2OBik-unfiltered-5-3	NA	<LOQ (0.0347)			<LOQ (0.00683)		
	MSU12129-H2OBik-unfiltered-5-4	NA	<LOQ (0.0347)			<LOQ (0.00683)		
	MSU12129-H2OBik-unfiltered-5-5	NA	E			E		
	MSU12129-H2OBik-unfiltered-5-6	NA	<LOQ (0.0696)			<LOQ (0.0171)		
	MSU12129-H2OBik-unfiltered-5-7	NA	<LOQ (0.0696)		NA	<LOQ (0.0171)		NA
	MSU12129-H2OBik-unfiltered-5-8	NA	<LOQ (0.0696)	<LOQ	NA	<LOQ (0.0171)	<LOQ	NA
Method Blk Filtered water	MSU12129-H2OBik-filtered-5-1	NA	NA			NA		
	MSU12129-H2OBik-filtered-5-2	NA	NA			NA		
	MSU12129-H2OBik-filtered-5-3	NA	E			E		
	MSU12129-H2OBik-filtered-5-4	NA	<LOQ (0.0347)			<LOQ (0.00683)		
	MSU12129-H2OBik-filtered-5-5	NA	<LOQ (0.0347)			<LOQ (0.00683)		
	MSU12129-H2OBik-filtered-5-6	NA	<LOQ (0.0696)			<LOQ (0.0171)		
	MSU12129-H2OBik-filtered-5-7	NA	<LOQ (0.0696)		NA	<LOQ (0.0171)		NA
	MSU12129-H2OBik-filtered-5-8	NA	<LOQ (0.0696)	<LOQ	NA	<LOQ (0.0171)	<LOQ	NA
Matrix Blk Fish Liver	FSH12129-LvrBlk-5-1	NA	0.0305			<LOQ (0.0171)		
	FSH12129-LvrBlk-5-2	NA	0.0331		32.1	<LOQ (0.0171)		NA
	FSH12129-LvrBlk-5-3	NA	0.0170	0.0269	0.00862	<LOQ (0.0171)	<LOQ	NA
Matrix Blk Rabbit Liver	RBL12129-LvrBlk-5-1	NA	<LOQ (0.0347)			<LOQ (0.00683)		
	RBL12129-LvrBlk-5-2	NA	<LOQ (0.0347)			<LOQ (0.00683)		
	RBL12129-LvrBlk-5-3	NA	<LOQ (0.0696)			<LOQ (0.0171)		
	RBL12129-LvrBlk-5-4	NA	<LOQ (0.0696)			<LOQ (0.0171)		
	RBL12129-LvrBlk-5-5*	NA	<LOQ (0.0696)		NA	<LOQ (0.0171)		NA
	RBL12129-LvrBlk-5-6*	NA	<LOQ (0.0696)	<LOQ	NA	<LOQ (0.0171)	<LOQ	NA
QC 250 ng/g	Mink Liver, D530, MS-5-1-1	NA	145%			61%		
	Mink Liver, D530, MSD-5-1-2*	NA	538%	342%	115%	50%	55%	21%
	CSL-1999030-03-01-MS-5-1-2*	NA	196%			116%		
	CSL-1999030-03-01-MSD-5-1-2*	NA	140%	168%	33%	112%	114%	4%
	LWL-1999029-12-MS-5-1-2*	NA	321%			57%		
	LWL-1999029-12-MSD-5-1-2*	NA	287%	304%	11%	68%	63%	18%
	BTL-1999040-01-MS-5-1-2*	NA	138%			81%		
	BTL-1999040-01-MSD-5-1-2*	NA	132%	135%	5%	84%	83%	4%
	TNL-TU54-MS-5-1-2	NA	68%			55%		
	TNL-TU54-MSD-5-1-2	NA	E	68%	NA	E	55%	NA
	FSL-P295-MS-5-1	NA	91%			97%		
	FSL-P295-MSD-5-1	NA	86%	88%	6%	84%	90%	15%
	PBL-980390LB-MS-5-1	NA	18%			70%		
	PBL-980390LB-MSD-5-2*	NA	179%	99%	163%	76%	73%	9%
	GFL-KZCKDM-D1-MS-5-1-2	NA	93%			82%		
	GFL-KZCKDM-D1-MSD-5-1	NA	105%	99%	12%	89%	86%	9%
	ITL-LCPT99503C-MS-5-1-2*	NA	90%			66%		
	ITL-LCPT99503C-MSD-5-1-2	NA	100%	95%	11%	72%	69%	8%
	MTL-10Vancleave98-MS-5-1-2	NA	79%			74%		
	MTL-10Vancleave98-MSD-5-1-2	NA	89%	84%	11%	84%	79%	13%
Liver Chinook Salmon	1999030-01	X	0.109			<LOQ (0.0171)		
	1999030-02	X	0.169			<LOQ (0.0171)		
	1999030-02-01	NA	0.0328			<LOQ (0.0171)		
	1999030-02-04	NA	0.126			<LOQ (0.0171)		
	1999030-03-01	NA	0.173		56.1	<LOQ (0.0171)		NA
	1999030-03-04	NA	0.0405	0.108	0.0608	<LOQ (0.0171)	<LOQ	NA
Liver Lake Whitefish	1999029-11	NA	0.0679			<LOQ (0.0171)		
	1999029-12	NA	0.0812			<LOQ (0.0171)		
	1999029-13	X	0.0738			<LOQ (0.0171)		
	1999029-14	X	0.0329		29.3	<LOQ (0.0171)		NA
	1999029-16	NA	0.0778	0.0667	0.0195	<LOQ (0.0171)	<LOQ	NA
Liver Brown Trout	1999040-01	NA	<LOQ (0.0174)			<LOQ (0.0171)		
	1999040-02	NA	<LOQ (0.0174)			<LOQ (0.0171)		
	1999040-03	NA	<LOQ (0.0174)			<LOQ (0.0171)		
	1999040-04	NA	<LOQ (0.0174)			<LOQ (0.0171)		
	1999040-05	NA	0.0255			<LOQ (0.0171)		
	1999040-06	NA	<LOQ (0.0174)			<LOQ (0.0171)		
	1999040-07	NA	<LOQ (0.0174)			<LOQ (0.0171)		
	1999040-08	NA	<LOQ (0.0174)			<LOQ (0.0171)		
	1999040-09	NA	<LOQ (0.0174)		NA	<LOQ (0.0171)		NA
	1999040-10	NA	<LOQ (0.0174)	<LOQ - 1 outlier	NA	<LOQ (0.0171)	<LOQ	NA

* High (>50%) surrogate deviations

Date Entered/Analyst: 12/22/99, 12/28/99, 12/29/99, 12/30/99,
01/12/00, 01/17/00, 01/18/00 LAC
Date Verified/Analyst: 0

NE = Not Extracted
E = Lost during extraction
NA = Not Applicable
LOQ = Limit of Quantitation
X = Verified PFOS concentration

PFOS = Perfluorooctanesulfonate
PFOSA = Perfluorooctane sulfonamide
PFHS = Perfluorohexanesulfonate
POAA = Perfluorooctanoate

FACT-GEN-030
MSU Environmental Samples

Study:
Product Number(Test Substance):
Matrix:
Method/Revision:
Analytical Equipment System Number:
Instrument Software/Version:
Date of Extraction/Analyst:
Date of Analysis/Analyst:
Date of Data Reduction/Analyst:

GEN030 MSU Environmental Samples
Various
Various
ETS-8-6.0 & ETS-8-7.0
Soup 020199, Amelia 062498
Masslynx 3.3
12/12/99 SAL/KJ/SRP/CSH
12/17/99, 12/20/99, 12/28/99, 12/29/99, 01/03/00, 01/06/00 IAS/MMH
12/20/99, 12/21/99, 12/22/99, 12/30/99, 01/03/00, 01/05/00, 01/07/00 MMH/IAS

Sample Data

Miscellaneous Liver

Group Dose	Sample #	Concentration of POAA ug/g or % Rec.	Mean POAA ug/g	RSD Std. Dev. MS/MSD RPD	Concentration of PFOSA ug/g or % Rec.	Mean PFOSA ug/g	RSD Std. Dev. MS/MSD RPD
Method Blk Unfiltered water	MSU12129-H2OBik-unfiltered-5-1	NA			NA		
	MSU12129-H2OBik-unfiltered-5-2	NA			NA		
	MSU12129-H2OBik-unfiltered-5-3	<LOQ (0.0719)			<LOQ (0.0188)		
	MSU12129-H2OBik-unfiltered-5-4	<LOQ (0.0719)			<LOQ (0.0188)		
	MSU12129-H2OBik-unfiltered-5-5	E			E		
	MSU12129-H2OBik-unfiltered-5-6	<LOQ (0.0719)			<LOQ (0.0188)		
	MSU12129-H2OBik-unfiltered-5-7	<LOQ (0.0719)		NA	<LOQ (0.0188)		NA
	MSU12129-H2OBik-unfiltered-5-8	<LOQ (0.0719)	<LOQ	NA	<LOQ (0.0188)	<LOQ	NA
Method Blk Filtered water	MSU12129-H2OBik-filtered-5-1	NA			NA		
	MSU12129-H2OBik-filtered-5-2	NA			NA		
	MSU12129-H2OBik-filtered-5-3	E			E		
	MSU12129-H2OBik-filtered-5-4	<LOQ (0.0719)			<LOQ (0.0188)		
	MSU12129-H2OBik-filtered-5-5	<LOQ (0.0719)			<LOQ (0.0188)		
	MSU12129-H2OBik-filtered-5-6	<LOQ (0.0719)			<LOQ (0.0188)		
	MSU12129-H2OBik-filtered-5-7	<LOQ (0.0719)		NA	<LOQ (0.0188)		NA
	MSU12129-H2OBik-filtered-5-8	<LOQ (0.0719)	<LOQ	NA	<LOQ (0.0188)	<LOQ	NA
Matrix Blk Fish Liver	FSH12129-LvrBlk-5-1	<LOQ (0.0719)			<LOQ (0.0188)		
	FSH12129-LvrBlk-5-2	<LOQ (0.0719)		NA	<LOQ (0.0188)		NA
	FSH12129-LvrBlk-5-3	<LOQ (0.0719)	<LOQ	NA	<LOQ (0.0188)	<LOQ	NA
Matrix Blk Rabbit Liver	RBL12129-LvrBlk-5-1	<LOQ (0.0719)			<LOQ (0.0188)		
	RBL12129-LvrBlk-5-2	<LOQ (0.0719)			<LOQ (0.0188)		
	RBL12129-LvrBlk-5-3	<LOQ (0.0719)			<LOQ (0.0188)		
	RBL12129-LvrBlk-5-4	<LOQ (0.0719)			<LOQ (0.0188)		
	RBL12129-LvrBlk-5-5*	<LOQ (0.0719)		NA	<LOQ (0.0188)		NA
	RBL12129-LvrBlk-5-6*	<LOQ (0.0719)	<LOQ	NA	<LOQ (0.0188)	<LOQ	NA
QC 250 ng/g	Mink Liver, D530, MS-S-1-1	86%			43%		
	Mink Liver, D530, MSD-S-1-2*	84%	85%	3%	81%	62%	60%
	CSL-1999030-03-01-MS-S-1-2*	131%			64%		
	CSL-1999030-03-01-MSD-S-1-2*	111%	121%	17%	52%	58%	21%
	LWL-1999029-12-MS-S-1-2*	133%			77%		
	LWL-1999029-12-MSD-S-1-2*	145%	139%	9%	83%	80%	7%
	BTL-1999040-01-MS-S-1-2*	127%			94%		
	BTL-1999040-01-MSD-S-1-2*	120%	124%	6%	92%	93%	3%
	TNL-TU54-MS-S-1-2	84%			71%		
	TNL-TU54-MSD-S-1-2	E	84%	NA	E	71%	NA
	FSL-P295-MS-S-1	72%			87%		
	FSL-P295-MSD-S-1	80%	76%	10%	82%	85%	7%
	PBL-980390LB-MS-S-1	85%			71%		
	PBL-980390LB-MSD-S-2*	125%	107%	41%	65%	68%	9%
	GFL-KZCKDM-D1-MS-S-1-2	82%			82%		
	GFL-KZCKDM-D1-MSD-S-1	103%	92%	23%	93%	88%	12%
	TTL-LCPT99503C-MS-S-1-2*	73%			78%		
	TTL-LCPT99503C-MSD-S-1-2	64%	69%	13%	71%	74%	10%
	MTL-10Vancleave98-MS-S-1-2	100%			74%		
	MTL-10Vancleave98-MSD-S-1-2	75%	88%	29%	74%	74%	0%
Liver Chinook Salmon	1999030-01	<LOQ (0.0719)			<LOQ (0.0188)		
	1999030-02	<LOQ (0.0719)			<LOQ (0.0188)		
	1999030-02-01	<LOQ (0.0719)			<LOQ (0.0188)		
	1999030-02-04	<LOQ (0.0719)			<LOQ (0.0188)		
	1999030-03-01	<LOQ (0.0719)		NA	<LOQ (0.0188)		NA
	1999030-03-04	<LOQ (0.0719)	<LOQ	NA	<LOQ (0.0188)	<LOQ	NA
Liver Lake Whitefish	1999029-11	<LOQ (0.0719)			<LOQ (0.0188)		
	1999029-12	<LOQ (0.0719)			<LOQ (0.0188)		
	1999029-13	<LOQ (0.0719)			<LOQ (0.0188)		
	1999029-14	<LOQ (0.0719)		NA	<LOQ (0.0188)		NA
	1999029-15	<LOQ (0.0719)			<LOQ (0.0188)		
	1999029-16	<LOQ (0.0719)	<LOQ	NA	<LOQ (0.0188)	<LOQ	NA
Liver Brown Trout	1999040-01	<LOQ (0.0719)			<LOQ (0.0188)		
	1999040-02	<LOQ (0.0719)			<LOQ (0.0188)		
	1999040-03	<LOQ (0.0719)			<LOQ (0.0188)		
	1999040-04	<LOQ (0.0719)			<LOQ (0.0188)		
	1999040-05	<LOQ (0.0719)			<LOQ (0.0188)		
	1999040-06	<LOQ (0.0719)			<LOQ (0.0188)		
	1999040-07	<LOQ (0.0719)			<LOQ (0.0188)		
	1999040-08	<LOQ (0.0719)			<LOQ (0.0188)		
	1999040-09	<LOQ (0.0719)		NA	<LOQ (0.0188)		NA
	1999040-10	<LOQ (0.0719)	<LOQ	NA	<LOQ (0.0188)	<LOQ	NA

* High (>50%) surrogate deviations

Date Entered/Analyst: 12/22/99, 12/28/99, 12/29/99, 12/30/99,
01/12/00, 01/17/00, 01/18/00 LAC
Date Verified/Analyst: 0

NB = Not Extracted
E = Lost during extraction
NA = Not Applicable
LOQ = Limit of Quantitation
X = Verified PFOS concentration

PFOS = Perfluorooctanesulfonate
PFOSA = Perfluorooctane sulfonamide
PFHS = Perfluorohexanesulfonate
POAA = Perfluorooctanoate

GEN-030 MSU Environmental Samples
Various
Various
ETS-4-6.0 & ETS-4-7.0
Samp 020199, Aerials 062498
Mandylux 3.3
12/17/99 SAL/KK/SRP/CSH
12/17/99, 12/20/99, 12/28/99, 12/29/99, 01/03/00, 01/05/00, 01/06/00, 01/08/00 LAS/NO/GH
12/28/99, 12/21/99, 12/22/99, 12/30/99, 01/03/00, 01/05/00, 01/06/00, 01/07/00, 01/11/00 MND/JOAS

Group Dose	Sample #	PFOS Verified	Concentration of PFOS ng/g or % Res.	Mean PFOS ng/g	RSD Std. Dev. MS/MSD RPD	Concentration of PFOS ng/g or % Res.	Mean PFOS ng/g	RSD Std. Dev. MS/MSD RPD
Method B1: Unfiltered water	MSU12129-HCOBB-#filtered-5-1	NA	NA			NA		
	MSU12129-HCOBB-#filtered-5-2	NA	NA			NA		
	MSU12129-HCOBB-#filtered-5-3	NA	<LOQ (0.0347)			<LOQ (0.00683)		
	MSU12129-HCOBB-#filtered-5-4	NA	<LOQ (0.0347)			<LOQ (0.00683)		
	MSU12129-HCOBB-#filtered-5-5	NA	E			E		
	MSU12129-HCOBB-#filtered-5-6	NA	<LOQ (0.0096)			<LOQ (0.0171)		
	MSU12129-HCOBB-#filtered-5-7	NA	<LOQ (0.0096)		NA	<LOQ (0.0171)		NA
	MSU12129-HCOBB-#filtered-5-8	NA	<LOQ (0.0096)	<LOQ		<LOQ (0.0171)	<LOQ	NA
Method B1: Filtered water	MSU12129-HCOBB-#filtered-5-1	NA	NA			NA		
	MSU12129-HCOBB-#filtered-5-2	NA	NA			NA		
	MSU12129-HCOBB-#filtered-5-3	NA	E			E		
	MSU12129-HCOBB-#filtered-5-4	NA	<LOQ (0.0347)			<LOQ (0.00683)		
	MSU12129-HCOBB-#filtered-5-5	NA	<LOQ (0.0347)			<LOQ (0.00683)		
	MSU12129-HCOBB-#filtered-5-6	NA	<LOQ (0.0096)			<LOQ (0.0171)		
	MSU12129-HCOBB-#filtered-5-7	NA	<LOQ (0.0096)		NA	<LOQ (0.0171)		NA
	MSU12129-HCOBB-#filtered-5-8	NA	<LOQ (0.0096)	<LOQ	NA	<LOQ (0.0171)	<LOQ	NA
Matrix B1: Fish Liver	FSH12129-L-wBB-5-1	NA	0.0303			<LOQ (0.0171)		
	FSH12129-L-wBB-5-2	NA	0.0331		32.1	<LOQ (0.0171)		NA
	FSH12129-L-wBB-5-3	NA	0.0170	0.0269	0.00842	<LOQ (0.0171)	<LOQ	NA
Matrix B1: Rabbit Liver	RBL12129-L-wBB-5-1	NA	<LOQ (0.0347)			<LOQ (0.00683)		
	RBL12129-L-wBB-5-2	NA	<LOQ (0.0347)			<LOQ (0.00683)		
	RBL12129-L-wBB-5-3	NA	<LOQ (0.0096)			<LOQ (0.0171)		
	RBL12129-L-wBB-5-4	NA	<LOQ (0.0096)			<LOQ (0.0171)		
	RBL12129-L-wBB-5-5*	NA	<LOQ (0.0096)		NA	<LOQ (0.0171)		NA
	RBL12129-L-wBB-5-6*	NA	<LOQ (0.0096)	<LOQ		<LOQ (0.0171)	<LOQ	NA
QC 250 ng/g	Mink Liver, D530, MS-5-1-1	NA	143%			81%		
	Mink Liver, D530, MS-5-1-2*	NA	538%	342%	115%	59%	55%	21%
	CSL-1999038-03-01-MS-5-1-2*	NA	258%			116%		
	CSL-1999038-03-01-MS-5-1-2*	NA	202%	230%	24%	112%	114%	4%
	LWL-1999028-12-MS-5-1-2*	NA	351%			57%		
	LWL-1999028-12-MS-5-1-2*	NA	317%	334%	10%	69%	67%	18%
	RTL-1999040-01-MS-5-1-2*	NA	138%			81%		
	RTL-1999040-01-MS-5-1-2*	NA	132%	135%	7%	86%	83%	4%
	TNL-TL04-MS-5-1-2	NA	68%			55%		
	TNL-TL04-MS-5-1-2	NA	E	68%	NA	E	55%	NA
	PSL-P295-MS-5-1	NA	111%			97%		
	PSL-P295-MS-5-1	NA	106%	108%	7%	84%	80%	15%
	PSL-98030LB-MS-5-1	NA	77%			70%		
	PSL-98030LB-MS-5-2*	NA	238%	158%	102%	70%	73%	9%
	GFL-KZK0DM-D1-MS-5-1-3	NA	101%			82%		
	GFL-KZK0DM-D1-MS-5-1-3	NA	112%	107%	11%	89%	86%	9%
	TTL-LCPT99903C-MS-5-1-2*	NA	92%			66%		
	TTL-LCPT99903C-MS-5-1-2	NA	102%	97%	10%	72%	69%	8%
	MTL-10V-mink-liver-MS-5-1-2	NA	108%			74%		
	MTL-10V-mink-liver-MS-5-1-2	NA	118%	113%	8%	84%	79%	15%
Liver Northern Fur Seal	PS3	NA	<LOQ (0.0347)			<LOQ (0.00683)		
	PS31*	X	0.122			<LOQ (0.00683)		
	PS3	NA	0.0547			<LOQ (0.00683)		
	97 CU 02	NA	NE			NE		
	98 CU KB 02	NA	<LOQ (0.0347)			<LOQ (0.00683)		

NE = Not Extracted
E = Lost during extraction
NA = Not Applicable
LOQ = Limit of Quantitation
X = Verified PFOS concentration

PFOH = Perfluorooctanoic acido
 PFOA = Perfluorooctano sulfoacido
 PFOS = Perfluorooctano sulfonato
 PFOA = Perfluorooctano

Study:
Product Number/Test Substance:
Matrix:
Method/Revision:
Analytical Equipment System Number:
Instrument Software/Version:
Date of Extraction/Analyst:
Date of Analysis/Analyst:
Date of Data Reduction/Analyst:
Sample Data

GENO 30 MSU Environmental Samples
Version
Version
ETS-6-6-0 & ETS-6-7-0
Scmp 020199, Amelia 062498
Mammya 3.3
121299 SAL/KC/SLP/CSH
121299, 123099, 122899, 122999, 010300, 010500, 010600, 010800 LAS/MNH
122599, 122199, 122299, 122699, 010400, 010500, 010600, 010700, 011000 MMT/IAS

Miscellaneous Liver

Group	Sample #	Concentration of PPOA µg/g or % Res.	Mean PPOA µg/g	RSD Std. Dev. MS/MSD RPD	Concentration of PPOA µg/g or % Res.	Mean PPOA µg/g	RSD Std. Dev. MS/MSD RPD
Method DB Unfiltered water	MSU12129-H2OBBR-unfiltered-5-1 MSU12129-H2OBBR-unfiltered-5-2 MSU12129-H2OBBR-unfiltered-5-3 MSU12129-H2OBBR-unfiltered-5-4 MSU12129-H2OBBR-unfiltered-5-5 MSU12129-H2OBBR-unfiltered-5-6 MSU12129-H2OBBR-unfiltered-5-7 MSU12129-H2OBBR-unfiltered-5-8	NA NA ≤LOQ (0.0719) ≤LOQ (0.0719) E ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719)	≤LOQ	NA NA	NA NA ≤LOQ (0.0188) ≤LOQ (0.0188) E ≤LOQ (0.0188) ≤LOQ (0.0188) ≤LOQ (0.0188)	≤LOQ	NA NA
Method DB Filtered water	MSU12129-H2OBBR-filtered-5-1 MSU12129-H2OBBR-filtered-5-2 MSU12129-H2OBBR-filtered-5-3 MSU12129-H2OBBR-filtered-5-4 MSU12129-H2OBBR-filtered-5-5 MSU12129-H2OBBR-filtered-5-6 MSU12129-H2OBBR-filtered-5-7 MSU12129-H2OBBR-filtered-5-8	NA NA E ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719)	≤LOQ	NA NA	NA NA E ≤LOQ (0.0188) ≤LOQ (0.0188) ≤LOQ (0.0188) ≤LOQ (0.0188) ≤LOQ (0.0188)	≤LOQ	NA NA
Matrix DB Fish Liver	FSH12129-LwBBR-5-1 FSH12129-LwBBR-5-2 FSH12129-LwBBR-5-3	≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719)	≤LOQ	NA NA	≤LOQ (0.0188) ≤LOQ (0.0188) ≤LOQ (0.0188)	≤LOQ	NA NA
Matrix DB Rabbit Liver	RBL12129-LwBBR-5-1 RBL12129-LwBBR-5-2 RBL12129-LwBBR-5-3 RBL12129-LwBBR-5-4 RBL12129-LwBBR-5-5 RBL12129-LwBBR-5-6	≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719)	≤LOQ	NA NA	≤LOQ (0.0188) ≤LOQ (0.0188) ≤LOQ (0.0188) ≤LOQ (0.0188) ≤LOQ (0.0188) ≤LOQ (0.0188)	≤LOQ	NA NA
QC 250 µg/g	Mink Liver, D530, MSB-5-1-1 Mink Liver, D530, MSD-5-1-2 CSL-1999030-03-01-MSB-5-1-2* CSL-1999030-03-01-MSD-5-1-2* LWL-1999028-12-MSB-5-1-2* LWL-1999028-12-MSD-5-1-2* BTL-1999040-01-MSB-5-1-2* BTL-1999040-01-MSD-5-1-2* TFL-TUSA-MSB-5-1-2 TFL-TUSA-MSD-5-1-2 PBL-P29-MSB-5-1 PBL-P29-MSD-5-1 PBL-98039LB-MSB-5-1 PBL-98039LB-MSD-5-2* OFL-KZC/KDM-D1-MSB-5-1-2 OFL-KZC/KDM-D1-MSD-5-1-2 TTL-LCPTIR9903C-MSB-5-1-2* TTL-LCPTIR9903C-MSD-5-1-2 MTL-10V-musclew98-MSB-5-1-2 MTL-10V-musclew98-MSD-5-1-2	86% 84% 131% 111% 133% 143% 127% 130% 84% E 72% 80% 13% 129% 82% 103% 73% 64% 100% 73%	83% 121% 17%	3% 17% 9% 6% NA NA 10% 41% 23% 13% NA NA	43% 81% 53% 41% 67% 73% 83% 81% 59% E 76% 71% 60% 54% 71% 82% 69% 39% 63% 63%	62% 47% 70% 82% 39% NA 73% 57% 77% 63% 63% ≤LOQ	69% 28% 8% 3% NA 8% 11% 14% 12% 1% NA NA
Liver Northern Fur Seal	P283 P283** P293 97 CU 02 98 CU KB 02 98 CU KB 03 98 CU KB 07 98 CU KB 09 98 CU KB 10 98 CU KB 11 98 CU KB 12 98 CU KB 13 98 CU KB 14 98 CU KB 15	≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) NE ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719)	≤LOQ	NA NA	≤LOQ (0.0188) ≤LOQ (0.0188) ≤LOQ (0.0188) NE ≤LOQ (0.0188) ≤LOQ (0.0188) ≤LOQ (0.0188) ≤LOQ (0.0188) ≤LOQ (0.0188) ≤LOQ (0.0188) ≤LOQ (0.0188) ≤LOQ (0.0188) ≤LOQ (0.0188) ≤LOQ (0.0188)	≤LOQ	NA NA
Liver Polar Bear	97D12 97D31 98D341 682-PLBR-0033 980127LB 980187LB 980290LB** 980543LB 980545LB 980112LB 980292LB 980294LB 980296LB 980603LB 980610LB 980633LB 980658LB	≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719)	≤LOQ	NA NA	≤LOQ (0.0188) ≤LOQ (0.0188) ≤LOQ (0.0188) ≤LOQ (0.0188) ≤LOQ (0.0188) ≤LOQ (0.0188) ≤LOQ (0.0188) ≤LOQ (0.0188) ≤LOQ (0.0188) ≤LOQ (0.0188) ≤LOQ (0.0188) ≤LOQ (0.0188) ≤LOQ (0.0188) ≤LOQ (0.0188) ≤LOQ (0.0188) ≤LOQ (0.0188) ≤LOQ (0.0188)	≤LOQ	NA NA
Liver Mink	D0530 D0566* D0590 D0618 D0630* D0684* D1000 D1024 D1050 D1092 D1110* D1134 D1190** D1194 D1196 D1244 D1248 D1660*	≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719) ≤LOQ (0.0719)	≤LOQ	NA NA	0.0334 0.0566 0.0209 0.0581 0.0703 0.0612 0.0294 ≤LOQ (0.0188) ≤LOQ (0.0188) ≤LOQ (0.0188) 0.0367 ≤LOQ (0.0188) 0.0340 0.0833 0.0742 ≤LOQ (0.0188) 0.0345 0.0621	0.0442 - 5 outliers	0.3 0.0201

* High (>50%) surrogate deviations confirmed.
 ** positive analyte confirmation was not achieved, used 499 -> 99 transition.
 Date Entered/Analyst: 12/12/99, 12/13/99, 12/29/99, 12/30/99,
 01/12/00, 01/17/00, 01/18/00, 01/19/00, 01/30/00 LAC
 Date Verified/Analyst: 0

NE = Not Extracted
E = Lost during extraction
NA = Not Applicable
LOQ = Limit of Quantitation
X = Verified PFOA concentration

PFOS = Perfluorooctanesulfonate
 PFOA = Perfluorooctanesulfonamid
 PFHxS = Perfluorohexanesulfonate
 POAA = Perfluorooctanoate

FACT-GEN-030
MSU Environmental Samples

Study:	GEN030 MSU Environmental Samples
Product Number(Test Substance):	Various
Matrix:	Various
Method/Revision:	ETS-8-6.0 & ETS-8-7.0
Analytical Equipment System Number:	Soup 020199, Amelia 061498
Instrument Software/Version:	Masslynx J.J
Date of Extraction/Analyst:	12/12/99 SAL/KK/SRP/CSH
Date of Analysis/Analyst:	12/17/99, 12/20/99, 12/28/99, 12/29/99, 01/03/00, 01/06/00, 02/01/00 IAS/MMH/MEE
Date of Data Reduction/Analyst:	12/20/99, 12/21/99, 12/22/99, 12/30/99, 01/03/00, 01/05/00, 01/07/00, 02/03/00 MMH/IAS/MEE

Miscellaneous Liver

Group Dose	Sample #	PFOS Verified	Concentration of PFOS µg/g or % Rec.	Mean PFOS µg/g	RSD Std. Dev. MS/MSD RPD	Concentration of PFHS µg/g or % Rec.	Mean PFHS µg/g	RSD Std. Dev. MS/MSD RPD
Method Blk Unfiltered water	MSU12129-H2OBK-unfiltered-5-1	NA	NA			NA		
	MSU12129-H2OBK-unfiltered-5-2	NA	NA			NA		
	MSU12129-H2OBK-unfiltered-5-3	NA	<LOQ (0.0347)			<LOQ (0.00683)		
	MSU12129-H2OBK-unfiltered-5-4	NA	<LOQ (0.0347)			<LOQ (0.00683)		
	MSU12129-H2OBK-unfiltered-5-5	NA	E			E		
	MSU12129-H2OBK-unfiltered-5-6	NA	<LOQ (0.0696)			<LOQ (0.0171)		
	MSU12129-H2OBK-unfiltered-5-7	NA	<LOQ (0.0696)		NA	<LOQ (0.0171)		NA
	MSU12129-H2OBK-unfiltered-5-8	NA	<LOQ (0.0696)	<LOQ	NA	<LOQ (0.0171)	<LOQ	NA
Method Blk Filtered water	MSU12129-H2OBK-filtered-5-1	NA	NA			NA		
	MSU12129-H2OBK-filtered-5-2	NA	NA			NA		
	MSU12129-H2OBK-filtered-5-3	NA	E			E		
	MSU12129-H2OBK-filtered-5-4	NA	<LOQ (0.0347)			<LOQ (0.00683)		
	MSU12129-H2OBK-filtered-5-5	NA	<LOQ (0.0347)			<LOQ (0.00683)		
	MSU12129-H2OBK-filtered-5-6	NA	<LOQ (0.0696)			<LOQ (0.0171)		
	MSU12129-H2OBK-filtered-5-7	NA	<LOQ (0.0696)		NA	<LOQ (0.0171)		NA
	MSU12129-H2OBK-filtered-5-8	NA	<LOQ (0.0696)	<LOQ	NA	<LOQ (0.0171)	<LOQ	NA
Matrix Blk Fish Liver	FSH12129-LvBk-5-1	NA	0.0305			<LOQ (0.0171)		
	FSH12129-LvBk-5-2	NA	0.0331		32.1	<LOQ (0.0171)		NA
	FSH12129-LvBk-5-3	NA	0.0170	0.0269	0.00861	<LOQ (0.0171)	<LOQ	NA
Matrix Blk Rabbit Liver	RBL12129-LvBk-5-1	NA	<LOQ (0.0347)			<LOQ (0.00683)		
	RBL12129-LvBk-5-2	NA	<LOQ (0.0347)			<LOQ (0.00683)		
	RBL12129-LvBk-5-3	NA	<LOQ (0.0696)			<LOQ (0.0171)		
	RBL12129-LvBk-5-4	NA	<LOQ (0.0696)			<LOQ (0.0171)		
	RBL12129-LvBk-5-5*	NA	<LOQ (0.0696)		NA	<LOQ (0.0171)		NA
	RBL12129-LvBk-5-6*	NA	<LOQ (0.0696)	<LOQ	NA	<LOQ (0.0171)	<LOQ	NA
QC 250 ng/g	Mink Liver, D530, MS-5-1-1	NA	145%			61%		
	Mink Liver, D530, MSD-5-1-2*	NA	538%	342%	115%	50%	55%	21%
	CSL-1999030-03-01-MS-5-1-2*	NA	196%			116%		
	CSL-1999030-03-01-MSD-5-1-2*	NA	140%	168%	33%	112%	114%	4%
	LWL-1999029-12-MS-5-1-2*	NA	321%			57%		
	LWL-1999029-12-MSD-5-1-2*	NA	287%	304%	11%	68%	63%	28%
	BTL-1999040-01-MS-5-1-2*	NA	138%			81%		
	BTL-1999040-01-MSD-5-1-2*	NA	132%	135%	5%	84%	83%	4%
	TNL-TU54-MS-5-1-2	NA	68%			55%		
	TNL-TU54-MSD-5-1-2	NA	E	68%	NA	E	55%	NA
	FSL-P295-MS-5-1	NA	91%			97%		
	FSL-P295-MSD-5-1	NA	86%	88%	6%	84%	90%	15%
	PBL-980390LB-MS-5-1	NA	18%			70%		
	PBL-980390LB-MSD-5-2*	NA	179%	99%	163%	76%	73%	9%
	GFL-KZCKDM-D1-MS-5-1-2	NA	93%			82%		
	GFL-KZCKDM-D1-MSD-5-1-1	NA	105%	99%	12%	89%	86%	9%
Liver Map Turtle	TTL-LCPT99503C-MS-5-1-2*	NA	90%			66%		
	TTL-LCPT99503C-MSD-5-1-2	NA	100%	95%	11%	72%	69%	8%
	MTL-10Vancleave98-MS-5-1-2	NA	79%			74%		
	MTL-10Vancleave98-MSD-5-1-2	NA	89%	84%	11%	84%	79%	13%
	F, #10, Vancleave 98	NA	0.0801			<LOQ (0.00683)		
	F, #09, Vancleave 98	NA	0.0514			<LOQ (0.00683)		
	F, #02, Lockville 98	NA	0.0739			<LOQ (0.00683)		
	F, #06, Lockville 99	NA	0.0394			<LOQ (0.00683)		
	M, (-1)	NA	0.703			<LOQ (0.00683)		
	F, (89, 8912)	NA	0.179			<LOQ (0.00683)		
Liver Terrapin	LCPT99503C	NA	<LOQ (0.0347)			<LOQ (0.00683)		
	LCPT99504C	NA	<LOQ (0.0347)		0.730	<LOQ (0.00683)		NA
	LCPT99505C	NA	<LOQ (0.0347)	0.188 - 3 outliers	0.257	<LOQ (0.00683)	<LOQ	NA
Liver Tuna	TU25*	X	<LOQ (0.0696)			<LOQ (0.0171)		
	TU34*	X	<LOQ (0.0696)			<LOQ (0.0171)		
	TU41*	X	<LOQ (0.0696)			<LOQ (0.0171)		
	TU48*	X	<LOQ (0.0696)			<LOQ (0.0171)		
	TU49	NA	<LOQ (0.0696)			<LOQ (0.0171)		
	TU54*	X	<LOQ (0.0696)			<LOQ (0.0171)		
	TU58	NA	<LOQ (0.0696)			<LOQ (0.0171)		
	TU83*	X	<LOQ (0.0696)			<LOQ (0.0171)		
	TU66	X	0.00698			<LOQ (0.0171)		
	TU84	NA	<LOQ (0.0696)			<LOQ (0.0171)		
	TU88	X	<LOQ (0.00696)		NA	<LOQ (0.0171)		NA
	TU90	NA	<LOQ (0.0696)	<LOQ - 1 outlier	NA	<LOQ (0.0171)	<LOQ	NA
Liver Green Frog	KZCKDM-D1	NA	<LOQ (0.0347)			<LOQ (0.00683)		
	KZCKDM-D2	X	0.285			<LOQ (0.00683)		
	Pool of 4 SJ0001	NA	<LOQ (0.0347)		NA	<LOQ (0.00683)		NA
		X	<LOQ (0.0347)	<LOQ - 1 outlier	NA	<LOQ (0.00683)	<LOQ	NA

* High (>50%) surrogate deviations

Date Entered/Analyst: 12/22/99, 12/28/99, 12/29/99, 12/30/99,
01/12/00, 01/17/00, 01/18/00, 02/04/00 LAC
Date Verified/Analyst:

NE = Not Extracted
E = Lost during extraction
NA = Not Applicable
LOQ = Limit of Quantitation
X = Verified PFOS concentration

PFOS = Perfluorooctanesulfonate
PFOSA = Perfluorooctane sulfonamide
PFHIS = Perfluorohexanesulfonate
POAA = Perfluorooctanoate

FACT-GEN-030
MSU Environmental Samples

Study:
Product Number(Test Substance):
Matrix:
Method/Revision:
Analytical Equipment System Number:
Instrument Software/Version:
Date of Extraction/Analyst:
Date of Analysis/Analyst:
Date of Data Reduction/Analyst:

GEN030 MSU Environmental Samples
Various
Various
ETS-8-6.0 & ETS-8-7.0
Soup 020199, Amelia 062498
Maatynex 3.3
12/12/99 SAL/KK/SRP/CSH
12/17/99, 12/20/99, 12/28/99, 12/29/99, 01/03/00, 01/06/00, 02/01/00 IAS/MMH/MEE
12/20/99, 12/21/99, 12/22/99, 12/30/99, 01/03/00, 01/05/00, 01/07/00, 02/03/00 MMH/IAS/MEE

Sample Data

Miscellaneous Liver

Group Dose	Sample #	Concentration of POAA ug/g or % Rec.	Mean POAA ug/g	RSD Std. Dev. MS/MSD RPD	Concentration of PFOSA ug/g or % Rec.	Mean PFOSA ug/g	RSD Std. Dev. MS/MSD RPD
Method Blk Unfiltered water	MSU12129-H2OBlk-unfiltered-5-1	NA			NA		
	MSU12129-H2OBlk-unfiltered-5-2	NA			NA		
	MSU12129-H2OBlk-unfiltered-5-3	<LOQ (0.0719)			<LOQ (0.0188)		
	MSU12129-H2OBlk-unfiltered-5-4	<LOQ (0.0719)			<LOQ (0.0188)		
	MSU12129-H2OBlk-unfiltered-5-5	E			E		
	MSU12129-H2OBlk-unfiltered-5-6	<LOQ (0.0719)			<LOQ (0.0188)		
	MSU12129-H2OBlk-unfiltered-5-7	<LOQ (0.0719)		NA	<LOQ (0.0188)		NA
	MSU12129-H2OBlk-unfiltered-5-8	<LOQ (0.0719)	<LOQ	NA	<LOQ (0.0188)	<LOQ	NA
Method Blk Filtered water	MSU12129-H2OBlk-filtered-5-1	NA			NA		
	MSU12129-H2OBlk-filtered-5-2	NA			NA		
	MSU12129-H2OBlk-filtered-5-3	E			E		
	MSU12129-H2OBlk-filtered-5-4	<LOQ (0.0719)			<LOQ (0.0188)		
	MSU12129-H2OBlk-filtered-5-5	<LOQ (0.0719)			<LOQ (0.0188)		
	MSU12129-H2OBlk-filtered-5-6	<LOQ (0.0719)			<LOQ (0.0188)		
	MSU12129-H2OBlk-filtered-5-7	<LOQ (0.0719)		NA	<LOQ (0.0188)		NA
	MSU12129-H2OBlk-filtered-5-8	<LOQ (0.0719)	<LOQ	NA	<LOQ (0.0188)	<LOQ	NA
Matrix Blk Fish Liver	FSH12129-LvrBlk-5-1	<LOQ (0.0719)			<LOQ (0.0188)		
	FSH12129-LvrBlk-5-2	<LOQ (0.0719)		NA	<LOQ (0.0188)		NA
	FSH12129-LvrBlk-5-3	<LOQ (0.0719)	<LOQ	NA	<LOQ (0.0188)	<LOQ	NA
Matrix Blk Rabbit Liver	RBL12129-LvrBlk-5-1	<LOQ (0.0719)			<LOQ (0.0188)		
	RBL12129-LvrBlk-5-2	<LOQ (0.0719)			<LOQ (0.0188)		
	RBL12129-LvrBlk-5-3	<LOQ (0.0719)			<LOQ (0.0188)		
	RBL12129-LvrBlk-5-4	<LOQ (0.0719)			<LOQ (0.0188)		
	RBL12129-LvrBlk-5-5*	<LOQ (0.0719)		NA	<LOQ (0.0188)		NA
	RBL12129-LvrBlk-5-6*	<LOQ (0.0719)	<LOQ	NA	<LOQ (0.0188)	<LOQ	NA
QC 250 ug/g	Mink Liver, D530, MS-5-1-1	84%			43%		
	Mink Liver, D530, MSD-5-1-2*	84%	85%	3%	81%	62%	60%
	CSL-1999030-03-01-MS-5-1-2*	131%			64%		
	CSL-1999030-03-01-MSD-5-1-2*	111%	121%	17%	52%	58%	21%
	LWL-1999029-12-MS-5-1-2*	133%			77%		
	LWL-1999029-12-MSD-5-1-2*	145%	139%	9%	83%	80%	7%
	BTL-1999040-01-MS-5-1-2*	127%			94%		
	BTL-1999040-01-MSD-5-1-2*	120%	124%	6%	92%	93%	3%
	TNL-TU54-MS-5-1-2	84%			71%		
	TNL-TU54-MSD-5-1-2	E	84%	NA	E	71%	NA
	PSL-P295-MS-5-1	72%			87%		
	PSL-P295-MSD-5-1	80%	76%	10%	82%	85%	7%
	PBL-980390LB-MS-5-1	85%			71%		
	PBL-980390LB-MSD-5-2*	129%	107%	41%	65%	68%	9%
	GFL-KZCKDM-D1-MS-5-1-2	82%			82%		
Liver Map Turtle	F, #10, Vanceave 98	<LOQ (0.0719)			<LOQ (0.0188)		
	F, #09, Vanceave 98	<LOQ (0.0719)			<LOQ (0.0188)		
	F, #02, Leeksville 98	<LOQ (0.0719)			<LOQ (0.0188)		
	F, #06, Leeksville 99	<LOQ (0.0719)			<LOQ (0.0188)		
	M, (-1)	<LOQ (0.0719)			<LOQ (0.0188)		
	F, (89, 8912)	<LOQ (0.0719)			<LOQ (0.0188)		
	LCPTR 9503C	<LOQ (0.0719)			<LOQ (0.0188)		
	LCPTR 9504C	<LOQ (0.0719)		NA	<LOQ (0.0188)		NA
	LCPTR 9505C	<LOQ (0.0719)	<LOQ	NA	<LOQ (0.0188)	<LOQ	NA
Liver Tuna	TU25*	<LOQ (0.0719)			<LOQ (0.0188)		
	TU34*	<LOQ (0.0719)			<LOQ (0.0188)		
	TU41*	<LOQ (0.0719)			<LOQ (0.0188)		
	TU48*	<LOQ (0.0719)			<LOQ (0.0188)		
	TU49	<LOQ (0.0719)			<LOQ (0.0188)		
	TU54*	<LOQ (0.0719)			<LOQ (0.0188)		
	TU58	<LOQ (0.0719)			<LOQ (0.0188)		
	TU63*	<LOQ (0.0719)			<LOQ (0.0188)		
	TU66	<LOQ (0.0719)			<LOQ (0.0188)		
	TU84	<LOQ (0.0719)		NA	<LOQ (0.0188)		NA
Liver Green Frog	TU88	<LOQ (0.0719)		NA	<LOQ (0.0188)		NA
	TU90	<LOQ (0.0719)	<LOQ	NA	<LOQ (0.0188)	<LOQ	NA
	KZCKDM-D1	<LOQ (0.0719)			<LOQ (0.0188)		
	KZCKDM-D2	<LOQ (0.0719)			<LOQ (0.0188)		
	Pool of 4	<LOQ (0.0719)		NA	<LOQ (0.0188)		NA
	SJ0001	<LOQ (0.0719)	<LOQ	NA	<LOQ (0.0188)	<LOQ	NA

* High (>50%) surrogate deviations

Date Entered/Analyst: 12/22/99, 12/23/99, 12/29/99, 12/30/99,
01/12/00, 01/17/00, 01/18/00, 02/04/00 LAC
Date Verified/Analyst:

NB = Not Extracted
E = Lost during extraction
NA = Not Applicable
LOQ = Limit of Quantitation
X = Verified PFOS concentration

PFOS = Perfluorooctanesulfonate
PFOSA = Perfluorooctane sulfonamide
PFHxS = Perfluorohexanesulfonate
POAA = Perfluorooctanoate

G

Study:
Product Number(Test Substance):
Matrix:
Method/Revision:
Analytical Equipment System Number:
Instrument Software/Version:
Date of Extraction/Analyst:
Date of Analysis/Analyst:
Date of Data Reduction/Analyst:
Sample Data

GEN033, MSU - Liver Samples

NA
Various livers - Unextracted Curves
BTS-8-6.0 and BTS-8-7.0
Devey 070799, Amelia 062498
Masslynx 3.3
03/14/00 SAL/CSH/KJK
03/16/00, 03/17/00, 3/19/00, 03/20/00, 03/21/00, 03/29/00, 04/07/00 IAS/MMH
03/20/00, 03/22/00, 03/23/00, 03/24/00, 04/04/00, 04/11/00 IAS/MMH

Filename: See Attachments
R-Squared Value: See Attachments
Slope: See Attachments
Y-Intercept: See Attachments

Various Livers

Group Dose	Sample #	Concentration of PFOS ug/g or % Rec	Mean PFOS ug/g	RSD Std. Dev. MS/MSD RPD	Concentration of POAA ug/g or % Rec	Mean POAA ug/g	RSD Std. Dev. MS/MSD RPD
Mink Liver	C23 (45)	0.501			< LOQ (0.0359 ug/g)		
	C26 (44)	0.443			< LOQ (0.0359 ug/g)		
	C27 (49)	0.145			< LOQ (0.0359 ug/g)		
	C33 (46)	0.435			< LOQ (0.0359 ug/g)		
	C35 (47)	0.355			< LOQ (0.0359 ug/g)		
	C37 (48)	0.833			< LOQ (0.0359 ug/g)		
	C42 (43)	0.420			< LOQ (0.0359 ug/g)		
	C44 (54)	0.237			< LOQ (0.0359 ug/g)		
	D10 (50)	1.67			< LOQ (0.0359 ug/g)		
	F15 (55)	0.548			< LOQ (0.0359 ug/g)		
	F19 (56)	0.783			< LOQ (0.0359 ug/g)		
	F21 (57)	1.03			< LOQ (0.0359 ug/g)		
	F24 (58)	0.868			< LOQ (0.0359 ug/g)		
	P01 (60)	2.16			< LOQ (0.0359 ug/g)		
	P03 (64)	4.80			< LOQ (0.0359 ug/g)		
	P09 (61)	0.841			< LOQ (0.0359 ug/g)		
	S11 (37)	0.902			< LOQ (0.0359 ug/g)		
	S15 (41)	1.27			< LOQ (0.0359 ug/g)		
	S18 (40)	1.99			< LOQ (0.0359 ug/g)		
	S19 (59)	2.68			< LOQ (0.0359 ug/g)		
	S25 (39)	0.509			< LOQ (0.0359 ug/g)		
	S30 (36)	0.184			< LOQ (0.0359 ug/g)		
	S35 (42)	0.0933			< LOQ (0.0359 ug/g)		
	S39 (38)	0.317			< LOQ (0.0359 ug/g)		
	T01 (51)	0.633			< LOQ (0.0359 ug/g)		
	T04 (53)	1.35			< LOQ (0.0359 ug/g)		
	T03 (52)	0.565			< LOQ (0.0359 ug/g)		
	V12 (62)	4.87			< LOQ (0.0359 ug/g)		
	V03 (65)	1.52		103	< LOQ (0.0359 ug/g)		NA
	V06 (63)	3.65	1.23	1.27	< LOQ (0.0359 ug/g)	< LOQ	NA
Baikal Seal Liver	J06 (81)	0.0127			< LOQ (0.0359 ug/g)		
	J09 (87)	0.0228			< LOQ (0.0718 ug/g)		
	J10 (86)	< LOQ (0.0347 ug/g)			< LOQ (0.0718 ug/g)		
	J12 (89)	0.0141			< LOQ (0.0359 ug/g)		
	J19 (84)	0.00931			< LOQ (0.0359 ug/g)		
	J20 (88)	0.0154			< LOQ (0.0718 ug/g)		
	J24 (82)	< LOQ (0.0347 ug/g)			< LOQ (0.0718 ug/g)		
	J27 (83)	< LOQ (0.0347 ug/g)			< LOQ (0.0718 ug/g)		
	J36 (85)	0.0146			< LOQ (0.0359 ug/g)		
	J37 (80)	< LOQ (0.0347 ug/g)			< LOQ (0.0718 ug/g)		
	R04 (69)	< LOQ (0.0347 ug/g)			< LOQ (0.0718 ug/g)		
	R13 (78)	0.0100			< LOQ (0.0359 ug/g)		
	R14 (74)	< LOQ (0.0347 ug/g)			< LOQ (0.0718 ug/g)		
	R16 (71)	0.00808			< LOQ (0.0359 ug/g)		
	R29 (72)	0.00795			< LOQ (0.0359 ug/g)		
	R42 (66)	0.0138			< LOQ (0.0359 ug/g)		
	R43 (73)	0.0156			< LOQ (0.0359 ug/g)		
	R45 (79)	0.00848			< LOQ (0.0359 ug/g)		
	R46 (77)	0.00778			< LOQ (0.0359 ug/g)		
	R47 (70)	< LOQ (0.0347 ug/g)			< LOQ (0.0718 ug/g)		
	R54 (67)	< LOQ (0.0347 ug/g)			< LOQ (0.0718 ug/g)		
	R55 (75)	0.0133			< LOQ (0.0718 ug/g)		
	R57 (76)	0.0158		33.7	< LOQ (0.0359 ug/g)		NA
	R64 (68)	0.00786	0.0123	0.00416	< LOQ (0.0359 ug/g)	< LOQ	NA
Common Dolphin Liver	L04 (91)	< LOQ (0.0347 ug/g)	NA	NA	< LOQ (0.0718 ug/g)	< LOQ	NA
	L05 (90)	0.0813	NA	NA	< LOQ (0.0718 ug/g)	< LOQ	NA

LOQ = Limit of Quantitation
NA = Not Applicable

PFOS = Perfluorooctanesulfonate
PFOSA = Perfluorooctanesulfonamide
PFHS = Perfluorohexanesulfonate
POAA = Perfluorooctanoic acid

Date Entered/Analyst: 03/28/00, 04/05/00, 04/07/00, 05/07/00 MN3/LAC
Date Verified/Analyst:

Study:
Product Number(Test Substance):
Matrix:
Method/Revision:
Analytical Equipment System Number:
Instrument Software/Version:
Date of Extraction/Analyst:
Date of Analysis/Analyst:
Date of Data Reduction/Analyst:
Sample Data

GEN033, MSU - Liver Samples
NA
Various livers - Unextracted Curves
ETS-6-6.0 and ETS-6-7.0
Devey 070799, Amelia 062498
Maalynx 3.3
03/14/00 SAL/CSH/KKK
03/16/00, 03/17/00, 3/19/00, 03/20/00, 03/21/00, 03/29/00, 04/07/00 IAS/MMH
03/20/00, 03/22/00, 03/23/00, 03/24/00, 04/04/00, 04/11/00 IAS/MMH

Filename: See Below
R-Squared Value: See Attachments
Slope: See Attachments
Y-Intercept: See Attachments

Various Livers

Group Dose	Sample #	Concentration of PFOSA ug/g or % Rec	Mean PFOSA ug/g	RSD Std. Dev. MS/MSD RPD	Concentration of PFHS ug/g or % Rec	Mean PFHS ug/g	RSD Std. Dev. MS/MSD RPD
Mink Liver	C23 (45)	0.0383			< LOQ (0.00643 ug/g)		
	C26 (44)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	C27 (49)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	C33 (46)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	C35 (47)	< LOQ (0.0376 ug/g)			0.00833		
	C37 (48)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	C42 (43)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	C44 (54)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	D10 (50)	0.0828			< LOQ (0.00643 ug/g)		
	F15 (55)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	F19 (56)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	F21 (57)	0.0579			< LOQ (0.00643 ug/g)		
	F24 (58)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	P01 (60)	0.551			< LOQ (0.00643 ug/g)		
	P03 (64)	0.590			0.0315		
	P09 (61)	< LOQ (0.0376 ug/g)			0.0852		
	S11 (57)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	S15 (41)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	S18 (40)	0.0414			< LOQ (0.00643 ug/g)		
	S19 (59)	0.132			0.0102		
	S25 (59)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	S30 (54)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	S35 (42)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	S39 (38)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	T01 (51)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	T04 (53)	< LOQ (0.0376 ug/g)			0.0104		
	T03 (52)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	V12 (62)	0.345			< LOQ (0.00643 ug/g)		
	V05 (65)	0.0594			< LOQ (0.00643 ug/g)		
	V08 (63)	0.0586	<LOQ - 10 Outliers	NA NA	< LOQ (0.00643 ug/g)	<LOQ - 5 Outliers	NA NA
Baikal Seal Liver	J06 (81)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	J09 (87)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	J10 (86)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	J12 (89)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	J19 (84)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	J20 (88)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	J24 (82)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	J27 (83)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	J36 (85)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	J37 (80)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	R04 (69)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	R13 (78)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	R14 (74)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	R16 (71)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	R29 (72)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	R42 (66)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	R43 (73)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	R45 (79)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	R46 (77)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	R47 (70)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	R54 (67)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	R55 (75)	< LOQ (0.0376 ug/g)			< LOQ (0.00643 ug/g)		
	R57 (76)	< LOQ (0.0376 ug/g)		NA	< LOQ (0.00643 ug/g)		NA
	R64 (68)	< LOQ (0.0376 ug/g)	<LOQ	NA	< LOQ (0.00643 ug/g)	<LOQ	NA
Ganges Dolphin Liver	L04 (91)	< LOQ (0.0376 ug/g)		NA	< LOQ (0.00643 ug/g)		NA
	L05 (90)	< LOQ (0.0376 ug/g)	<LOQ	NA	< LOQ (0.00643 ug/g)	<LOQ	NA

LOQ = Limit of Quantitation
NA = Not Applicable

PFOS = Perfluorooctanesulfonate
PFOSA = Perfluorooctanesulfonamide
PFHS = Perfluorohexane sulfonate
POAA = Perfluorooctanoate

Date Entered/Analyst: 03/22/00, 04/05/00, 04/07/00, 05/07/00 IAS/MMH/LAC
Date Verified/Analyst:

Study:
Product Number/Test Substance:
Matrix:
Method/Revision:
Analytical Equipment System Number:
Instrument Software/Version:
Date of Extraction/Analysis:
Date of Analysis/Analysis:
Date of Data Reduction/Analysis:
Sample Data

GEN-033, MSU - Liver Samples

NA

Various livers - unextracted curves

ETS-6-6.0 and ETS-6-7.0

Dewey 070799, Amelco 062498

Model 3.3

03/14/00 SAL/CSH/KKK

03/16/00, 03/17/00, 3/19/00, 03/20/00, 03/21/00, 03/29/00, 04/07/00 IAS/MMH

03/20/00, 03/22/00, 03/23/00, 03/24/00, 04/04/00, 04/11/00 IAS/MMH

Filename: See Attachments

R-Squared Value: See Attachments

Slope: See Attachments

Y-intercept: See Attachments

Various Livers

Group Date	Sample #	Concentration of PFOS ng/g or % Rec	Mean PFOS ng/g	RSD Std. Dev. MS/MSD RPD	Concentration of FOAA ng/g or % Rec	Mean FOAA ng/g	RSD Std. Dev. MS/MSD RPD
Common Liver Female, Adult	6P (1)F.A	0.0432			0.0414		
	29F (5)F.A**	0.0565			0.102		
	30F (6)F.A	0.0483			0.0506		
	32F (7)F.A	0.0977			0.0897		
	34F (9)F.A**	0.0335		61.9	0.0297		56.9
	42F (11)F.A	0.150	0.0715	0.0442	0.143	0.0761	0.0432
Common Liver Female, Juvenile	7F (5)F.J	0.0913			0.0841		
	9F (5)F.J	0.468		113	0.444		115
	34F (10)F.J	0.0499	0.203	0.230	0.0467	0.192	0.219
Common Liver Male, Juvenile	22F (6)M.J	0.0433			0.0454		
	33F (8)M.J	0.0316		17.2	0.0303		25.7
	44F (12)M.J	0.0337	0.0362	0.00623	0.0299	0.0350	0.00899
Bottlenose Dolphin Liver	T6P1 (13)	0.181			<LOQ (0.0359 ng/g)		
	T6P1 (17)	0.296			<LOQ (0.0718 ng/g)		
	T6P1 (14)	0.169			<LOQ (0.0718 ng/g)		
	T6A11 (14)	0.425		44.5	<LOQ (0.0718 ng/g)		NA
	T6K12 (15)	<LOQ (0.00896 ng/g)	0.268 - One Outlier	0.119	<LOQ (0.0718 ng/g)	<LOQ	NA
Striped Dolphin Liver	SCQ3 (18)	0.161			<LOQ (0.0718 ng/g)		
	SCV1 (19)	0.0891			<LOQ (0.0718 ng/g)		
	SCF03 (20)	0.0944		40.2	<LOQ (0.0718 ng/g)		NA
	SCF04 (21)	0.0647	0.102	0.0410	<LOQ (0.0718 ng/g)	<LOQ	NA
Weddell Seal Liver	W31 (23)	<LOQ (0.0347 ng/g)	<LOQ	NA	<LOQ (0.0718 ng/g)	<LOQ	NA
Swordfish Liver	S23 (23)	<LOQ (0.00896 ng/g)			<LOQ (0.0359 ng/g)		
	S24 (24)	<LOQ (0.00896 ng/g)			<LOQ (0.0359 ng/g)		
	S25 (25)	0.00774			<LOQ (0.0359 ng/g)		
	S32 (27)	<LOQ (0.00896 ng/g)		NA	<LOQ (0.0359 ng/g)		NA
	S48 (28)	0.0173	<LOQ - Two Outliers	NA	<LOQ (0.0359 ng/g)	<LOQ	NA
Tussock Liver	T1 (25)	<LOQ (0.0347 ng/g)			<LOQ (0.0359 ng/g)		
	T1L17.56 (25)	0.0433			<LOQ (0.0718 ng/g)		
	T2 (29)	0.0674			<LOQ (0.0359 ng/g)		
	T15 (28)	0.0568			<LOQ (0.0718 ng/g)		
	T17 (31)	0.0491			<LOQ (0.0359 ng/g)		
	T20 (32)	0.0207			<LOQ (0.0359 ng/g)		
	T23 (30)	0.0560		46.2	<LOQ (0.0359 ng/g)		NA
	T25 (34)	0.0259	0.0483 - One Outlier	0.0223	<LOQ (0.0718 ng/g)	<LOQ	NA
Blacktail Gull Liver	BHC01 (100)	0.292			<LOQ (0.0359 ng/g)		
	BHC02 (101)	0.260			<LOQ (0.0359 ng/g)		
	BHC03 (102)	0.148			<LOQ (0.0359 ng/g)		
	BHC04 (103)	0.593			<LOQ (0.0359 ng/g)		
	BHC05 (104)	0.271			<LOQ (0.0359 ng/g)		
	BTCH005 (92)	0.0881			<LOQ (0.0359 ng/g)		
	BTCH006 (99)	0.107			<LOQ (0.0359 ng/g)		
	BTCH010 (96)	0.215			<LOQ (0.0359 ng/g)		
	BTCH011 (98)	0.143			<LOQ (0.0359 ng/g)		
	BTCH012 (97)	0.0705			<LOQ (0.0359 ng/g)		
	BTCH0401 (90)	0.126			<LOQ (0.0359 ng/g)		
	BTCH0401 (94)	0.0737			<LOQ (0.0359 ng/g)		
	BTCH0401 (95)	0.0707			<LOQ (0.0359 ng/g)		
	HRC04 (106)	0.116		69.2	<LOQ (0.0359 ng/g)		NA
	HRC09 (105)	0.0939	0.172	0.119	<LOQ (0.0359 ng/g)	<LOQ	NA

** NO PFOS confirmation performed.

LOQ = Limit of Quantitation

NA = Not Applicable

Date Entered/Analyst:

03/28/00, 04/05/00, 04/07/00, 05/07/00 MGH/LAC

Date Verified/Analyst:

PFOS = Perfluorooctanesulfonate

FOAA = Perfluorooctanoate

PFOSA = Perfluorooctanesulfonamide

PFHS = Perfluorohexanesulfonate

Study:
Product Number/Test Substance):
Matrix:
Method/Revision:
Analytical Equipment System Number:
Instrument Software/Version:
Date of Extraction/Analyte:
Date of Analysis/Analyte:
Date of Data Reduction/Analyte:
Sample Data

GEN-033, NSU - Liver Samples

NA
Various livers - unextracted curves
STS-4-6.0 and STS-4-7.0
Davy 070799, Amels 042498
Masslynx 3.3

Filename: See Below
R-Squared Value: See Attachments
Slope: See Attachments
Y-Intercept: See Attachments

03/16/00, 03/17/00, 3/19/00, 03/20/00, 03/21/00, 03/29/00, 04/07/00 IAS/MMH
03/20/00, 03/22/00, 03/23/00, 03/24/00, 04/04/00, 04/11/00 IAS/MMH

Various Livers

Group Date	Sample #	Concentration of PFOS ug/g or % Res	Mean PFOS ug/g	RSD Std. Dev. MS/MSD RPD	Concentration of PFOS ug/g or % Res	Mean PFOS ug/g	RSD Std. Dev. MS/MSD RPD
Common Liver Female, Adult	6F (1)F.A	<LOQ (0.0376 ug/g)			<LOQ (0.0063 ug/g)		
	39F (5)F.A**	<LOQ (0.0376 ug/g)			<LOQ (0.0063 ug/g)		
	30F (6)F.A	<LOQ (0.0376 ug/g)			<LOQ (0.0063 ug/g)		
	32F (7)F.A	<LOQ (0.0376 ug/g)			<LOQ (0.0063 ug/g)		
	34F (9)F.A**	<LOQ (0.0376 ug/g)			<LOQ (0.0063 ug/g)		NA
	42F (11)F.A	<LOQ (0.0376 ug/g)	<LOQ	NA	<LOQ (0.0063 ug/g)	<LOQ	NA
Common Liver Female, Juvenile	7F (3)F.J	0.0888			<LOQ (0.0063 ug/g)		
	9F (5)F.J	<LOQ (0.0376 ug/g)		NA	<LOQ (0.0063 ug/g)		NA
	36F (10)F.J	<LOQ (0.0376 ug/g)	<LOQ - One Outlier	NA	<LOQ (0.0063 ug/g)	<LOQ	NA
Common Liver Male, Juvenile	22F (4)M.J	<LOQ (0.0376 ug/g)		NA	<LOQ (0.0063 ug/g)		
	33F (8)M.J	<LOQ (0.0376 ug/g)		NA	<LOQ (0.0063 ug/g)		NA
	64F (12)M.J	<LOQ (0.0376 ug/g)	<LOQ	NA	<LOQ (0.0063 ug/g)	<LOQ	NA
Bottlenose Dolphin Liver	T0591 (13)	0.224			<LOQ (0.0063 ug/g)		
	T0691 (17)	0.358			<LOQ (0.0063 ug/g)		
	T0611 (14)	0.129		54.1	<LOQ (0.0063 ug/g)		
	T0611 (16)	0.129		9.103	<LOQ (0.0063 ug/g)		
	T0612 (15)	0.115	9.191		<LOQ (0.0063 ug/g)	<LOQ	NA
Striped Dolphin Liver	SCF02 (16)	<LOQ (0.0376 ug/g)			<LOQ (0.0063 ug/g)		
	SCF71 (19)	<LOQ (0.0376 ug/g)			<LOQ (0.0063 ug/g)		
	SCF03 (20)	<LOQ (0.0376 ug/g)		NA	<LOQ (0.0063 ug/g)		NA
	SCF04 (21)	<LOQ (0.0376 ug/g)	<LOQ - One Outlier	NA	<LOQ (0.0063 ug/g)	<LOQ - One Outlier	NA
Wedded Seal Liver	W31 (22)	<LOQ (0.0376 ug/g)	<LOQ	NA	<LOQ (0.0063 ug/g)	<LOQ	NA
Burreedfish Liver	S23 (23)	<LOQ (0.0376 ug/g)			0.0954		
	S24 (24)	<LOQ (0.0376 ug/g)			<LOQ (0.0063 ug/g)		
	S25 (25)	<LOQ (0.0376 ug/g)			<LOQ (0.0063 ug/g)		
	S33 (27)	<LOQ (0.0376 ug/g)		NA	<LOQ (0.0063 ug/g)		NA
	S48 (26)	<LOQ (0.0376 ug/g)	<LOQ	NA	<LOQ (0.0063 ug/g)	<LOQ - One Outlier	NA
Tussock Liver	T1 (33)	<LOQ (0.0376 ug/g)			<LOQ (0.0063 ug/g)		
	T1L7156 (35)	<LOQ (0.0376 ug/g)			<LOQ (0.0063 ug/g)		
	T2 (29)	<LOQ (0.0376 ug/g)			<LOQ (0.0063 ug/g)		
	T15 (28)	<LOQ (0.0376 ug/g)			<LOQ (0.0063 ug/g)		
	T17 (21)	<LOQ (0.0376 ug/g)			<LOQ (0.0063 ug/g)		
	T20 (22)	<LOQ (0.0376 ug/g)			<LOQ (0.0063 ug/g)		
	T23 (20)	<LOQ (0.0376 ug/g)		NA	<LOQ (0.0063 ug/g)		NA
	T25 (24)	<LOQ (0.0376 ug/g)	<LOQ	NA	<LOQ (0.0063 ug/g)	<LOQ	NA
	BH001 (100)	<LOQ (0.0376 ug/g)			<LOQ (0.0063 ug/g)		
	BH002 (101)	<LOQ (0.0376 ug/g)			<LOQ (0.0063 ug/g)		
Blacktail Gull Liver	BH003 (102)	<LOQ (0.0376 ug/g)			<LOQ (0.0063 ug/g)		
	BH004 (103)	<LOQ (0.0376 ug/g)			<LOQ (0.0063 ug/g)		
	BH005 (104)	<LOQ (0.0376 ug/g)			<LOQ (0.0063 ug/g)		
	BTC0303 (92)	<LOQ (0.0376 ug/g)			<LOQ (0.0063 ug/g)		
	BTC0304 (99)	<LOQ (0.0376 ug/g)			<LOQ (0.0063 ug/g)		
	BTC0510 (96)	<LOQ (0.0376 ug/g)			<LOQ (0.0063 ug/g)		
	BTC0511 (98)	<LOQ (0.0376 ug/g)			<LOQ (0.0063 ug/g)		
	BTC0512 (97)	<LOQ (0.0376 ug/g)			<LOQ (0.0063 ug/g)		
	BTC0601 (93)	<LOQ (0.0376 ug/g)			<LOQ (0.0063 ug/g)		
	BTC0602 (94)	<LOQ (0.0376 ug/g)			<LOQ (0.0063 ug/g)		
	BTC0603 (95)	<LOQ (0.0376 ug/g)			<LOQ (0.0063 ug/g)		
	BH004 (106)	<LOQ (0.0376 ug/g)		NA	<LOQ (0.0063 ug/g)		NA
	BH009 (105)	<LOQ (0.0376 ug/g)	<LOQ	NA	<LOQ (0.0063 ug/g)	<LOQ	NA

** NO PFOS confirmation performed.

LOQ = Limit of Quantitation

NA = Not Applicable

Date Recd/Analyte:
Date Verified/Analyte:

03/28/00, 04/05/00, 04/07/00, 05/07/00 IAS/MMH

PFOS = Perfluorooctanesulfonate

POAA = Perfluorooctanesulfonamide

PFOSA = Perfluorooctanesulfonamide

PFHS = Perfluorohexanesulfonate

Study:
Product Number(Test Substance):
Matrix:
Method/Revision:
Analytical Equipment System Number:
Instrument Software/Version:
Date of Extraction/Analysis:
Date of Analysis/Analyst:
Date of Data Reduction/Analyst:
Sample Data

GEN033, MSU - Liver Samples
NA
Various livers - unextracted curves
ETS-8-6.0 and ETS-4-7.0
Dewey 070799, Amelia 061498
Mamlynx 3.3
05/14/00 SAL/CSH/KKK
05/16/00, 05/17/00, 3/19/00, 03/20/00, 03/21/00, 03/29/00, 04/07/00 LAS/MCH
05/29/00, 05/22/00, 05/23/00, 05/24/00, 04/04/00, 04/11/00 LAS/MCH

Filename: See Attachments
R-Squared Value: See Attachments
Slope: See Attachments
Y-Intercept: See Attachments

Various Livers

Group Date	Sample #	Concentration of PFOS ng/g or % Rec	Mean PFOS ng/g	RSD Std. Dev. MS/MSD RPD	Concentration of POAA ng/g or % Rec	Mean POAA ng/g	RSD Std. Dev. MS/MSD RPD
Method Bk	RBL03140-H2O Bk-S-1	< LOQ (0.00696 ng/g)			< LOQ (0.0359 ng/g)		
	RBL03140-H2O Bk-S-2	< LOQ (0.00696 ng/g)			< LOQ (0.0359 ng/g)		
	RBL03140-H2O Bk-S-3	< LOQ (0.00696 ng/g)			< LOQ (0.0718 ng/g)		
	RBL03140-H2O Bk-S-4	< LOQ (0.00696 ng/g)			< LOQ (0.0718 ng/g)		
	RBL03140-H2O Bk-S-5	< LOQ (0.00696 ng/g)			< LOQ (0.0718 ng/g)		
	RBL03140-H2O Bk-S-6	< LOQ (0.0247 ng/g)			< LOQ (0.00719 ng/g)		
	RBL03140-H2O Bk-S-7	< LOQ (0.00696 ng/g)			< LOQ (0.0718 ng/g)		
	RBL03140-H2O Bk-S-8	< LOQ (0.00696 ng/g)			< LOQ (0.0718 ng/g)		
	RBL03140-H2O Bk-S-9	< LOQ (0.00696 ng/g)			< LOQ (0.0718 ng/g)		
	RBL03140-H2O Bk-S-10	< LOQ (0.0247 ng/g)			< LOQ (0.00719 ng/g)		
	RBL03140-H2O Bk-S-11	< LOQ (0.0247 ng/g)			< LOQ (0.0359 ng/g)		
	RBL03140-H2O Bk-S-12	< LOQ (0.0247 ng/g)		NA	< LOQ (0.0359 ng/g)		NA
	RBL03140-H2O Bk-S-13	< LOQ (0.0247 ng/g)	< LOQ	NA	< LOQ (0.0359 ng/g)	< LOQ	NA
Matrix Bk	RBL03140-Liver Bk-S-1	< LOQ (0.00696 ng/g)			< LOQ (0.0718 ng/g)		
	RBL03140-Liver Bk-S-2	< LOQ (0.00696 ng/g)			< LOQ (0.0718 ng/g)		
	RBL03140-Liver Bk-S-3	< LOQ (0.00696 ng/g)			< LOQ (0.0718 ng/g)		
	RBL03140-Liver Bk-S-4	< LOQ (0.00696 ng/g)			< LOQ (0.0718 ng/g)		
	RBL03140-Liver Bk-S-5	0.00883			< LOQ (0.0359 ng/g)		
	RBL03140-Liver Bk-S-6	< LOQ (0.00696 ng/g)			< LOQ (0.0718 ng/g)		
	RBL03140-Liver Bk-S-7	< LOQ (0.00696 ng/g)			< LOQ (0.0718 ng/g)		
	RBL03140-Liver Bk-S-8	< LOQ (0.00696 ng/g)			< LOQ (0.0718 ng/g)		
	RBL03140-Liver Bk-S-9	< LOQ (0.00696 ng/g)			< LOQ (0.0718 ng/g)		
	RBL03140-Liver Bk-S-10	< LOQ (0.00696 ng/g)			< LOQ (0.0718 ng/g)		
	RBL03140-Liver Bk-S-11	< LOQ (0.0247 ng/g)		NA	< LOQ (0.0359 ng/g)		NA
	RBL03140-Liver Bk-S-12	< LOQ (0.0247 ng/g)		NA	< LOQ (0.0359 ng/g)		NA
	RBL03140-Liver Bk-S-13	< LOQ (0.0247 ng/g)	< LOQ - One Outlier	NA	< LOQ (0.0359 ng/g)	< LOQ	NA
QC 250 ppb	RBL03140-MSD-1 ppb-S-1*	104%			112%		
	RBL03140-MSD-1 ppb-S-2*	114%	109%	9%	106%	109%	8%
	RBL03140-MSD-250 ppb-S-1	120%			128%		
	RBL03140-MSD-250 ppb-S-2	132%	127%	10%	128%	128%	0%
	44F (12)-MS	98%			106%		
	44F (12)-MSD	107%	103%	9%	109%	107%	2%
	T08/91 (17)-MS	131%			59%		
	T08/91 (17)-MSD	140%	133%	6%	64%	61%	9%
	SCV1 (19)-MS	66%			56%		
	SCV1 (19)-MSD	93%	79%	35%	76%	66%	30%
	WS1 (22)-MS	81%			63%		
	WS1 (22)-MSD	73%	77%	16%	62%	62%	2%
	324 (24)-MSD	92%	NA	NA	101%	NA	NA
	T15 (26)-MS	84%			93%		
	T15 (26)-MSD	103%	94%	21%	105%	99%	12%
	T11/F156 (35)-MS	93%			92%		
	T11/F156 (35)-MSD	72%	82%	25%	80%	86%	15%
	S19 (39)-MS	126%			126%		
	S19 (39)-MSD	234%	280%	33%	154%	140%	30%
	P9 (61)-MS	211%			102%		
	P9 (61)-MSD	142%	187%	48%	112%	107%	9%
	R47 (70)-MS	68%			70%		
	R47 (70)-MSD	69%	69%	2%	68%	69%	3%
	L4 (91)-MS	69%			73%		
	L4 (91)-MSD	187%	128%	92%	165%	119%	78%
	BTG0302 (92)-MS	143%	NA	NA	155%	NA	NA
	BTG0302 (92)-MSD	122%			137%		
	BTG0302 (94)-MSD	126%	121%	2%	156%	146%	13%

LOQ = Limit of Quantitation
NA = Not Applicable

PFOS = Perfluorooctanesulfonate
POAA = Perfluorooctanoate
PFOSA = Perfluorooctanesulfonamide
PFHS = Perfluorohexanesulfonate

Date Entered/Analyst: 03/22/00, 04/05/00, 04/07/00, 05/07/00 MCH/VLAC
Date Verified/Analyst:

Study:
Product Number(Test Substance):
Matrix:
Method/Revision:
Analytical Equipment System Number:
Instrument Software/Version:
Date of Analysis/Analyst:
Date of Data Reduction/Analyst:
Sample Data

GEN013, MSU - Liver Samples
NA
Various livers - unextracted curves
ETS-4-6.0 and ETS-4-7.0
Devery 070799, Amelia 062498
Masslynx 3.3
03/16/00, 03/17/00, 3/19/00, 03/20/00, 03/21/00, 03/29/00, 04/07/00 IAS/MMH
03/20/00, 03/22/00, 03/23/00, 03/24/00, 04/04/00, 04/11/00 IAS/MMH

Filename: See Below
R-Squared Value: See Attachments
Slope: See Attachments
Y-Intercept: See Attachments

Various Livers

Group Dose	Sample #	Concentration of PFOSA ng/g or % Rec	Mean PFOSA ng/g	RSD Std. Dev. MS/MSD RPD	Concentration of PFHS ng/g or % Rec	Mean PFHS ng/g	RSD Std. Dev. MS/MSD RPD
Method Blk	RBL03140-H20 Blk-5-1	< LOQ (0.0376 ng/g)			<LOQ (0.00683 ng/g)		
	RBL03140-H20 Blk-5-2	< LOQ (0.0376 ng/g)			<LOQ (0.00683 ng/g)		
	RBL03140-H20 Blk-5-3	< LOQ (0.0376 ng/g)			<LOQ (0.00683 ng/g)		
	RBL03140-H20 Blk-5-4	< LOQ (0.0376 ng/g)			<LOQ (0.00683 ng/g)		
	RBL03140-H20 Blk-5-5	< LOQ (0.0376 ng/g)			<LOQ (0.00683 ng/g)		
	RBL03140-H20 Blk-5-6	< LOQ (0.0376 ng/g)			<LOQ (0.00683 ng/g)		
	RBL03140-H20 Blk-5-7	< LOQ (0.0376 ng/g)			<LOQ (0.00683 ng/g)		
	RBL03140-H20 Blk-5-8	< LOQ (0.0376 ng/g)			<LOQ (0.00683 ng/g)		
	RBL03140-H20 Blk-5-9	< LOQ (0.0376 ng/g)			<LOQ (0.00683 ng/g)		
	RBL03140-H20 Blk-5-10	< LOQ (0.0376 ng/g)			<LOQ (0.00683 ng/g)		
	RBL03140-H20 Blk-5-11	< LOQ (0.0376 ng/g)			<LOQ (0.00683 ng/g)		
	RBL03140-H20 Blk-5-12	< LOQ (0.0376 ng/g)		NA	<LOQ (0.00683 ng/g)		NA
	RBL03140-H20 Blk-5-13	< LOQ (0.0376 ng/g)	<LOQ	NA	<LOQ (0.00683 ng/g)	<LOQ	NA
Matrix Blk	RBL03140-Liver Blk-5-1	< LOQ (0.0376 ng/g)			<LOQ (0.00683 ng/g)		
	RBL03140-Liver Blk-5-2	< LOQ (0.0376 ng/g)			<LOQ (0.00683 ng/g)		
	RBL03140-Liver Blk-5-3	< LOQ (0.0376 ng/g)			<LOQ (0.00683 ng/g)		
	RBL03140-Liver Blk-5-4	< LOQ (0.0376 ng/g)			<LOQ (0.00683 ng/g)		
	RBL03140-Liver Blk-5-5	< LOQ (0.0376 ng/g)			<LOQ (0.00683 ng/g)		
	RBL03140-Liver Blk-5-6	< LOQ (0.0376 ng/g)			<LOQ (0.00683 ng/g)		
	RBL03140-Liver Blk-5-7	< LOQ (0.0376 ng/g)			<LOQ (0.00683 ng/g)		
	RBL03140-Liver Blk-5-8	< LOQ (0.0376 ng/g)			<LOQ (0.00683 ng/g)		
	RBL03140-Liver Blk-5-9	< LOQ (0.0376 ng/g)			<LOQ (0.00683 ng/g)		
	RBL03140-Liver Blk-5-10	< LOQ (0.0376 ng/g)			<LOQ (0.00683 ng/g)		
	RBL03140-Liver Blk-5-11	< LOQ (0.0376 ng/g)		NA	<LOQ (0.00683 ng/g)		NA
	RBL03140-Liver Blk-5-12	< LOQ (0.0376 ng/g)		NA	<LOQ (0.00683 ng/g)		NA
	RBL03140-Liver Blk-5-13	< LOQ (0.0376 ng/g)	<LOQ	NA	<LOQ (0.00683 ng/g)	<LOQ	NA
QC 250 ppb	RBL03140-MSD-1 ppb-5-1*	67%			87%		
	RBL03140-MSD-1 ppb-5-2*	67%	67%	2%	57%	72%	42%
	RBL03140-MSD-250 ppb-5-1	114%			61%		
	RBL03140-MSD-250 ppb-5-2	84%	99%	30%	48%	54%	25%
	44P (12)-MSD	94%			84%		
	44P (12)-MSD	89%	92%	3%	87%	88%	3%
	T18/91 (17)-MSD	148%			63%		
	T18/91 (17)-MSD	191%	168%	28%	63%	63%	1%
	SCV1 (19)-MSD	53%			54%		
	SCV1 (19)-MSD	87%	70%	48%	64%	59%	17%
	WS1 (22)-MSD	69%			55%		
	WS1 (22)-MSD	69%	69%	0%	60%	60%	1%
	S24 (24)-MSD	98%	NA	NA	76%	NA	NA
	T15 (28)-MSD	68%			67%		
	T15 (28)-MSD	95%	81%	36%	102%	84%	40%
	T1LP156 (35)-MSD	86%			88%		
	T1LP156 (35)-MSD	66%	76%	36%	73%	80%	19%
	S19 (39)-MSD	140%			97%		
	S19 (39)-MSD	2%	71%	195%	-1%	48%	203%
	P9 (61)-MSD	73%			52%		
	P9 (61)-MSD	68%	70%	7%	48%	50%	6%
	R47 (70)-MSD	68%			55%		
	R47 (70)-MSD	78%	73%	14%	58%	57%	5%
	L4 (91)-MSD	56%			54%		
	L4 (91)-MSD	119%	85%	82%	145%	102%	86%
	BTG0902 (92)-MSD	70%	NA	NA	31%	NA	NA
	BTG0902 (92)-MSD	63%			18%		
	BTG0902 (92)-MSD	63%	63%	2%	22%	20%	20%

LOQ = Limit of Quantitation
NA = Not Applicable

PFOS = Perfluorooctanesulfonate
FOAA = Perfluorooctanoate
PFOSA = Perfluorooctanesulfonamide
PFHS = Perfluorohexanesulfonate

Date Entered/Analyst: 03/28/00, 04/05/00, 04/07/00, 05/07/00 bmg/vlac
Date Verified/Analyst:

3M ENVIRONMENTAL LABORATORY

METHOD

EXTRACTION OF POTASSIUM PERFLUOROOCTANESULFONATE OR OTHER FLUOROchemical COMPOUNDS FROM SERUM FOR ANALYSIS USING HPLC- ELECTROSPRAY/MASS SPECTROMETRY

Method Number: ETS-8-4.1

Adoption Date: 03/01/99

Revision Date:

Author: Lisa Clemen, Glenn Langenburg

Approved By:

Laboratory Manager

Date

Group Leader

Date

Technical Reviewer

Date

1.0 SCOPE AND APPLICATION

- 1.1 Scope:** This method is for the extraction of potassium perfluorooctanesulfonate (PFOS) or other fluorochemical compounds from serum.
- 1.2 Applicable compounds:** Fluorochemical surfactants or other fluorinated compounds.
- 1.3 Matrices:** Rabbit, rat, bovine, monkey, and human serum or other fluids as designated in the validation report.

2.0 SUMMARY OF METHOD

- 2.1 This method describes the procedure for extracting potassium perfluorooctanesulfonate (PFOS) or other fluorochemical surfactants from serum, or other fluids, using an ion pairing reagent and methyl-*tert*-butyl ether (MtBE). In this method, seven fluorochemicals were extracted: PFOS, PFOSA, PFOSAA, EtFOSE-OH, PFOSEA, M556, and surrogate standard (see 3.0 *Definitions*). An ion pairing reagent is added to the sample and the analyte ion pair is partitioned into MtBE. The MtBE extract is removed and put onto a nitrogen evaporator until dry. Each extract is reconstituted in 1.0 mL of methanol, then filtered through a 3 cc plastic syringe attached to a 0.2 μ m nylon filter into glass autovials.
- 2.2 These sample extracts are analyzed following method ETS-8-5.1 or other appropriate method.

3.0 DEFINITIONS

- 3.1 PFOS: perfluorooctanesulfonate (anion of potassium salt) $C_8F_{17}SO_3^-$
- 3.2 PFOSA: perfluorooctane sulfonylamide $C_8F_{17}SO_2NH_2$
- 3.3 PFOSAA: perfluorooctane sulfonylamido (ethyl)acetate $C_8F_{17}SO_2N(CH_2CH_3)CH_2CO_2^-$
- 3.4 EtFOSE-OH: 2(N-ethylperfluorooctane sulfonamido)-ethyl alcohol $C_8F_{17}SO_2N(CH_2CH_3)CH_2CH_2OH$
- 3.5 PFOSEA: perfluorooctane sulfonyl ethylamide $C_8F_{17}SO_2N(CH_2CH_3)H$
- 3.6 M556: $C_8F_{17}SO_2N(H)(CH_2COOH)$
- 3.7 Surrogate standard: 1H-1H-2H-2H perfluorooctane sulfonic acid

4.0 WARNINGS AND CAUTIONS

4.1 Health and safety warnings

- 4.1.1 Use universal precautions, especially laboratory coats, goggles, and gloves when handling animal tissue, which may contain pathogens.

5.0 INTERFERENCES

- 5.1 There are no interferences known at this time.

6.0 EQUIPMENT

- 6.1 The following equipment is used while performing this method. Equivalent equipment is acceptable.
- 6.1.1 Vortex mixer, VWR, Vortex Genie 2
- 6.1.2 Centrifuge, Mistral 1000 or IEC
- 6.1.3 Shaker, Eberbach or VWR
- 6.1.4 Nitrogen evaporator, Organomation

6.1.5 Balance (± 0.100 g)

7.0 SUPPLIES AND MATERIALS

- 7.1 Gloves
- 7.2 Eppendorf or disposable pipettes
- 7.3 Nalgene bottles, capable of holding 250 mL and 1 L
- 7.4 Volumetric flasks, glass, type A
- 7.5 I-CHEM vials, glass, 40 mL glass
- 7.6 Centrifuge tubes, polypropylene, 15 mL
- 7.7 Labels
- 7.8 Oxford Dispenser –3.0 to 10.0 mL
- 7.9 Syringes, capable of measuring 5 μ L to 50 μ L
- 7.10 Graduated pipettes
- 7.11 Syringes, disposable plastic, 3 cc
- 7.12 Syringe filters, nylon, 0.2 μ m, 25 mm
- 7.13 Timer
- 7.14 Crimp cap autovials and caps
- 7.15 Crimpers

Note: Prior to using glassware and bottles, rinse 3 times with methanol and 3 times with Milli-Q™ water. Rinse syringes a minimum of 9 times with methanol, 3 rinses from 3 separate vials.

8.0 REAGENTS AND STANDARDS

- 8.1 Type I reagent grade water, Milli-Q™ or equivalent; all water used in this method should be Milli-Q™ water and may be provided by a Milli-Q TOC Plus™ system
- 8.2 Sodium hydroxide (NaOH), J.T Baker or equivalent
- 8.3 Tetrabutylammonium hydrogen sulfate(TBA), Kodak or equivalent
- 8.4 Sodium carbonate (Na_2CO_3), J.T. Baker or equivalent
- 8.5 Sodium bicarbonate (NaHCO_3), J.T. Baker or equivalent
- 8.6 Methyl-T-Butyl Ether, Omnisolv, glass distilled or HPLC grade
- 8.7 Methanol, Omnisolv, glass distilled or HPLC grade
- 8.8 Serum or blood, frozen from supplier
- 8.9 **Fluorochemical standards**
 - 8.9.1 PFOS (3M Specialty Chemical Division), molecular weight = 538
 - 8.9.2 PFOSA (3M Specialty Chemical Division), molecular weight = 499
 - 8.9.3 PFOSAA (3M Specialty Chemical Division), molecular weight = 585

- 8.9.4 EtFOSE-OH (3M Specialty Chemical Division), molecular weight = 570
- 8.9.5 PFOSEA (3M Specialty Chemical Division), molecular weight = 527
- 8.9.6 M556 (3M Specialty Chemical Division), molecular weight = 557
- 8.9.7 Surrogate standard: 4-H, perfluorooctane sulfonic acid (1-H, 1-H, 2-H, 2-H $C_8F_{13}SO_3H$) molecular weight = 428
- 8.9.8 Other fluorochemicals, as appropriate

8.10 Reagent preparation

NOTE: When preparing larger volumes than listed in reagent, standard, or surrogate preparation, adjust accordingly.

- 8.10.1 10 N sodium hydroxide (NaOH): Weigh approximately 200 g NaOH. Pour into a 1000 mL beaker containing 500 mL Milli-Q™ water, mix until all solids are dissolved. Store in a 1 L Nalgene bottle.
- 8.10.2 1 N sodium hydroxide (NaOH): Dilute 10 N NaOH 1:10. Measure 10 mL of 10 N NaOH solution into a 100 mL volumetric flask and dilute to volume using Milli-Q™ water. Store in a 125 mL Nalgene bottle.
- 8.10.3 0.5 M tetrabutylammonium hydrogen sulfate (TBA): Weigh approximately 169 g of TBA into a 1 L volumetric containing 500 mL Milli-Q™ water. Adjust to pH 10 using approximately 44 to 54 mL of 10 N NaOH (While adding the last mL of NaOH, add slowly because the pH changes abruptly). Dilute to volume with Milli-Q™ water. Store in a 1 L Nalgene bottle.
 - 8.10.3.1 TBA requires a check prior to each use to ensure pH = 10. Adjust as needed using 1 N NaOH solution.
- 8.10.4 0.25 M sodium carbonate/sodium bicarbonate buffer ($Na_2CO_3/NaHCO_3$): Weigh approximately 26.5 g of sodium carbonate (Na_2CO_3) and 21.0 g of sodium bicarbonate ($NaHCO_3$) into a 1 L volumetric flask and bring to volume with Milli-Q™ water. Store in a 1 L Nalgene bottle.

8.11 Standards preparation

- 8.11.1 Prepare PFOS standards for the standard curve.
- 8.11.2 Prepare other fluorochemical standards, as appropriate. Multicomponent fluorochemical standards are acceptable (for example, one working standard solution containing 1.00 ppm PFOS, 1.02 ppm PFOSA, 0.987 ppm PFOSAA, and 1.10 ppm EtFOSE-OH.)
- 8.11.3 Weigh approximately 100 mg of PFOS into a 100 mL volumetric flask and record the actual weight.
- 8.11.4 Bring to volume with methanol for a stock standard of approximately 1000 ppm ($\mu g/mL$).
- 8.11.5 Dilute the stock solution with methanol for a working standard 1 solution of approximately 50 ppm.
- 8.11.6 Dilute working standard 1 with methanol for a working standard 2 solution of approx. 5.0 ppm.

8.11.7 Dilute working standard 1 with methanol for a working standard 3 solution of approx. 0.50 ppm.

8.12 Surrogate stock standard preparation

8.12.1 Weigh approximately 50-60 mg of surrogate standard 1-H, 1-H, 2-H, 2-H, $C_8F_{13}SO_3H$ into a 50 mL volumetric flask and record the actual weight.

8.12.2 Bring to volume with methanol for a surrogate stock of approximately 1000-1200 ppm.

8.12.3 Prepare a surrogate working standard. Transfer approximately 1 mL of surrogate stock to a 10 mL volumetric flask and bring to volume with methanol for a working standard of 100 ppm. Record the actual volume transferred.

9.0 SAMPLE HANDLING

9.1 All samples are received frozen and must be kept frozen until the extraction is performed.

9.2 Allow samples to thaw to room temperature prior to extraction.

10.0 QUALITY CONTROL

10.1 Solvent Blanks, Method blanks and matrix blanks

10.1.1 An aliquot of 1.0 mL methanol is used as a solvent blank.

10.1.2 Extract two 1.0 mL aliquots of Milli-QTM water following this procedure and use as method blanks.

10.1.3 Extract two 1.0 mL aliquots of the serum following this procedure and use as matrix blanks. See 11.1.4.

10.2 Matrix spikes

10.2.1 Prepare and analyze matrix spike and matrix spike duplicate samples to determine the accuracy of the extraction.

10.2.2 Prepare each spike using a sample chosen by the analyst, usually the control matrix received with each sample set.

10.2.3 Expected concentrations will fall in the mid-range of the initial calibration curve. Additional spikes may be included and may fall in the low-range of the initial calibration curve.

10.2.4 Prepare one matrix spike and matrix spike duplicate per 40 samples, with a minimum of 2 matrix spikes per batch.

10.3 Continuing calibration checks

10.3.1 Prepare continuing calibration check samples to ensure the accuracy of the initial calibration curve.

10.3.2 Prepare, at a minimum, one continuing check per group of 10 samples. For example, if a sample set = 34, four checks are prepared and extracted.

10.3.3 Prepare each continuing calibration check from the same matrix used to prepare the initial curve.

- 10.3.4** The expected concentrations will fall within the mid-range of the initial calibration curve. Additional spikes may be included that fall in the low-range of the initial calibration curve. This is necessary if the analyst must quantitate using only the low end of the calibration curve (for example, 5 ppb – 100 ppb, rather than 5 ppb – 1000 ppb).

11.0 CALIBRATION AND STANDARDIZATION

11.1 Prepare matrix calibration standards

- 11.1.1** Transfer 1 mL of serum to a 15 mL centrifuge tube.
- 11.1.2** If most sample volumes are less than 1.0 mL, extract standards with matrix volumes equal to the sample volumes. Do not extract less than 0.50 mL of matrix. Record each sample volume on the extraction sheet.
- 11.1.3** While preparing a total of twenty aliquots in 15 mL centrifuge tubes, mix or shake between aliquots.
- 11.1.4** Two 1 mL aliquots, or other appropriate volume, serve as matrix blanks. Typically use the standard concentrations and spiking amounts listed in Table 1, at the end of this section, to spike, in duplicate, two standard curves, for a total of eighteen standards, two matrix blanks, and two method blanks.
- 11.1.5** Refer to validation report ETS-8-4.0 & ETS-8-5.0-V-1, which lists the working ranges and the Linear Calibration Range (LCR) for calibration curves.
- 11.1.6** Use Attachment D as an aid in calculating the concentrations of the working standards. See Section 13.0 to calculate actual concentrations of PFOS in calibration standards.
- 11.2** To each standard, blank, or continuing check, add appropriate amount of surrogate working standard for the concentration to fall within the calibration curve range 5 ppb - 1000 ppb.
- 11.3** Extract spiked matrix standards following 12.6-12.16 of this method. Use these standards to establish each initial curve on the mass spectrometer.

Table 1 Approximate spiking amounts for standards and spikes Using 1.0 mL of matrix		
Working standard (approx. conc.)	μL	Approx. final conc. of analyte in matrix
-	-	Blank
0.500 ppm	10	0.005 ppm
0.500 ppm	20	0.010 ppm
5.00 ppm	5	0.025 ppm
5.00 ppm	10	0.050 ppm
5.00 ppm	20	0.100 ppm
50.0 ppm	5	0.250 ppm
50.0 ppm	10	0.500 ppm
50.0 ppm	15	0.750 ppm
50.0 ppm	20	1.00 ppm

12.0 PROCEDURE

- 12.1 Obtain frozen samples and allow to thaw at room temperature or in a lukewarm waterbath.
- 12.2 Vortex mix for 15 seconds, then transfer 1.0 mL or other appropriate volume to a 15 mL polypropylene centrifuge tube.
- 12.3 Return unused samples to freezer after extraction amounts have been removed.
- 12.4 Record the initial volume on the extraction worksheet.
- 12.5 Label the tube with the study number, sample ID, date and analyst initials. See attached worksheet for documenting the remaining steps.
- 12.6 Spike all samples, including blanks and standards, ready for extraction with surrogate standard as described in 11.2.
- 12.7 Spike each matrix with the appropriate amount of standard as described in 11.1, or Table 1 in that section, for the calibration curve standards. Also prepare matrix spikes and continuing calibration standards.
- 12.8 Vortex mix the standard curve samples, matrix spike samples, and continuing calibration samples for 15 seconds.
- 12.9 Check to ensure the 0.5 M TBA reagent is at pH 10. If not, adjust accordingly.
- 12.10 To each sample, add 1 mL 0.5 M TBA and 2 mL of 0.25M sodium carbonate/sodium bicarbonate buffer.
- 12.11 Using an Oxford Dispenser, add 5 mL methyl-*tert*-butyl ether.
- 12.12 Cap each sample and put on the shaker at a setting of 300 rpm, for 20 minutes.
- 12.13 Centrifuge for 20 to 25 minutes at a setting of 3500 rpm, or until layers are well separated.

- 12.14 Label a fresh 15 mL centrifuge tube with the same information as in 12.5.
- 12.15 Remove 4.0 mL of the organic layer to this clean 15 mL centrifuge tube.
- 12.16 Put each sample on the analytical nitrogen evaporator until dry, approximately 1 to 2 hours.
- 12.17 Add 1.0 mL of methanol to each centrifuge tube using a graduated pipette.
- 12.18 Vortex mix for 30 seconds.
- 12.19 Attach a 0.2 µm nylon mesh filter to a 3 cc syringe and transfer the sample to this syringe. Filter into a 1.5 mL glass autovial or low-volume autovial when necessary.
- 12.20 Label the autovial with the study number, animal number and gender, sample timepoint, matrix, final solvent, extraction date, and analyst(s) performing the extraction.
- 12.21 Cap and store extracts at room temperature or at approximately 4 °C until analysis.
- 12.22 Complete the extraction worksheet, attached to this document, and tape in the study notebook or include in study binder, as appropriate.

13.0 DATA ANALYSIS AND CALCULATIONS

13.1 Calculations

- 13.1.1 Calculate actual concentrations of PFOS, or other applicable fluorochemical, in calibration standards using the following equation:

$$\frac{\text{mL of standard} \times \text{concentration of standard } (\mu\text{g / mL})}{\text{mL of standard} + \text{mL of surrogate standard} + \text{initial matrix volume (mL)}} =$$

Final Concentration (µg/mL) of PFOS in matrix

14.0 METHOD PERFORMANCE

- 14.1 The method detection limit (MDL) is analyte and matrix specific. Refer to MDL report for specific MDL and limit of quantitation (LOQ) values (see **Attachments B and C**).
- 14.2 The following quality control samples are extracted with each batch of samples to evaluate the quality of the extraction and analysis.
 - 14.2.1 Method blanks and matrix blanks.
 - 14.2.2 Matrix spike and matrix spike duplicate samples to determine accuracy and precision of the extraction.
 - 14.2.3 Continuing calibration check samples to determine the continued accuracy of the initial calibration curve.
- 14.3 Refer to section 14 of ETS-8-5.1 for method performance criteria.

15.0 POLLUTION PREVENTION AND WASTE MANAGEMENT

- 15.1 Sample waste is disposed in biohazard containers, flammable solvent waste is disposed in high BTU containers, and used glass pipette waste is disposed in broken glass containers located in the laboratory.

16.0 RECORDS

- 16.1 Complete the extraction worksheet attached to this method, and tape in the study notebook or include in the 3-ring study binder, as appropriate.

17.0 ATTACHMENTS

- 17.1 Attachment A, Extraction worksheet
17.2 Attachment B, MDL/LOQ values and summary
17.3 Attachment C, Calibration standard concentration worksheet

18.0 REFERENCES

- 18.1 The validation report associated with this method is ETS-8-4.0 & 5.0-V-1.
18.2 FACT-M-3.1, "Analysis of Serum or Other Fluid Extracts for Fluorochemicals using HPLC-Electrospray Mass Spectrometry"

19.0 AFFECTED DOCUMENTS

- 19.1 ETS-8-5.1, "Analysis of Serum or Other Fluid Extracts for Fluorochemicals using HPLC-Electrospray Mass Spectrometry"

20.0 REVISIONS

<u>Revision Number</u>	<u>Reason For Revision</u>	<u>Revision Date</u>
1	Section 12.21 Changed to include sample storage at room temperature. Section 12.13 Added the shaker speed. Section 12.17 Final volume is 1.0 mL; not adjusted for initial volumes less than 1.0 mL.	04/02/99

- 4.1.2 When handling samples or solvents wear appropriate protective gloves, eyewear, and clothing.

4.2 Cautions:

- 4.2.1 Operate the solvent pumps below a back pressure of 400 bar (5800 psi). If the back pressure exceeds 400 bar, the HP1100 will initiate automatic shutdown.
- 4.2.2 Do not run solvent pumps to dryness.

5.0 INTERFERENCES

- 5.1 To minimize interferences when analyzing samples, Teflon shall not be used for sample storage or any part of instrumentation that comes in contact with the sample or extract.

6.0 EQUIPMENT

- 6.1 Equipment listed below may be modified in order to optimize the system. Document any modifications in the raw data as method deviations.
 - 6.1.1 Micromass Quattro II triple quadrupole Mass Spectrometer equipped with an electrospray ionization source.
 - 6.1.2 HP1100 low pulse solvent pumping system, solvent degasser, column compartment, and autosampler

7.0 SUPPLIES AND MATERIALS

- 7.1 Supplies
 - 7.1.1 High purity grade air regulated to approximately 100 psi (house air system)
 - 7.1.2 HPLC analytical column, specifics to be determined by the analyst and documented in the raw data
 - 7.1.3 Capped autovials or capped 15 ml centrifuge tubes

8.0 REAGENTS AND STANDARDS

- 8.1 Reagents
 - 8.1.1 Methanol, HPLC grade or equivalent
 - 8.1.2 Milli-Q™ water (ASTM type I), all water used in this method should be ATSM type I, or equivalent, and be provided by a Milli-Q TOC Plus system or other vendor
 - 8.1.3 Ammonium acetate, reagent grade or equivalent
 - 8.1.3.1 When preparing different amounts than those listed, adjust accordingly.
 - 8.1.3.2 2.0 mM ammonium acetate solution: Weigh approximately 0.300 g ammonium acetate. Pour into a 2000 mL volumetric container containing 2000 mL Milli-Q™ water, mix until all solids are dissolved. Store at room temperature.

8.2 Standards

- 8.2.1** Typically two method blanks, two matrix blanks, and eighteen matrix standards are prepared during the extraction procedure. Refer to ETS-8-6.0.

9.0 SAMPLE HANDLING

- 9.1** Fresh matrix standards are prepared with each analysis. Extracted standards and samples are stored in capped autovials or capped 15 ml centrifuge tubes until analysis.
- 9.2** If analysis will be delayed, extracted standards and samples may be stored at room temperature, or refrigerated at approximately 4° C, until analysis can be performed.

10.0 QUALITY CONTROL

10.1 Method Blanks and Matrix Blanks

- 10.1.1** Solvent blanks, method blanks, and matrix blanks are prepared and analyzed with each batch to determine contamination or carryover.
- 10.1.2** Analyze a method blank and a matrix blank prior to each calibration curve.

10.2 Matrix Spikes

- 10.2.1** Matrix spikes are prepared and analyzed to determine the matrix effect on the recovery efficiency.
- 10.2.2** Matrix spike duplicates are prepared and analyzed to measure the precision and the recovery for each analyte.
- 10.2.3** Analyze a matrix spike and matrix spike duplicate per forty samples. With a minimum of 2 spikes per batch.
- 10.2.4** Matrix spike and matrix spike duplicate concentrations will fall in the mid-range of the initial calibration curve. Additional spike concentrations may fall in the low-range of the initial calibration curve.

10.3 Continuing Calibration Checks

- 10.3.1** Continuing calibration verifications are analyzed to verify the continued accuracy of the calibration curve.
- 10.3.2** Analyze a mid-range calibration standard every tenth sample, with a minimum of one per batch.

11.0 CALIBRATION AND STANDARDIZATION

- 11.1** Analyze the extracted matrix standards prior to and following each set of sample extracts. The average of two standard curves will be plotted by linear regression ($y = mx + b$), weighted $1/x$, not forced through the origin, using MassLynx or other suitable software.
- 11.2** If the curve does not meet requirements perform routine maintenance or reextract the standard curve (if necessary) and reanalyze.

- 11.3** For purposes of accuracy when quantitating low levels of analyte, it may be necessary to use the low end of the calibration curve rather than the full range of the standard curve. Example: when attempting to quantitate approximately 10 ppb of analyte, generate a calibration curve consisting of the standards from 5 ppb to 100 ppb rather than the full range of the curve (5 ppb to 1000 ppb). This will reduce inaccuracy attributed to linear regression weighting of high concentration standards.

12.0 PROCEDURES

12.1 Acquisition Set up

12.1.1 Set up the sample list.

12.1.1.1 Assign a sample list filename using MO-DAY-last digit of year-increasing letter of the alphabet starting with a

12.1.1.2 Assign a method (MS file) for acquiring

12.1.1.3 Assign an HPLC program (Inlet file)

12.1.1.4 Type in sample descriptions and vial position numbers

12.1.2 To create a method click on method in the Acquisition control panel then mass spectrometer headings and select SIR (Single Ion Recording) or MRM (Multiple Reaction Monitoring). Set Ionization Mode as appropriate and mass to 499 or other appropriate masses. A full scan is usually collected along with the SIRs. Save acquisition method. If MS/MS instruments are employed, additional product ion fragmentation information may be collected. Refer to Micromass MassLynx GUIDE TO DATA ACQUISITION for additional information and MRM.

12.1.3 Typically the analytical batch run sequence begins and ends with a set of extracted matrix standards.

12.1.4 Samples are analyzed with a continuing calibration verification injected standard after every tenth sample. Solvent blanks should be analyzed periodically to monitor possible analyte carryover and are not considered samples but may be included as such.

12.2 Using the Autosampler

12.2.1 Set up sample tray according to the sample list prepared in Section 12.1.1.

12.2.2 Set-up the HP1100/autosampler at the following conditions or at conditions the analyst considers appropriate for optimal response. Record actual conditions in the instrument logbook:

12.2.2.1 Sample size = 10 μ L injection

12.2.2.2 Inject/sample = 1

12.2.2.3 Cycle time = 9 minutes

12.2.2.4 Solvent ramp conditions

Time	MeOH	2.0 mM Ammonium acetate
0.00 min.	40%	60%
1.0 min.	40%	60%
4.5 min.	95%	5%
6.5 min.	95%	5%
7.0 min.	40%	60%
9.0 min.	40%	60%

12.2.2.5 Press the "Start" button.

12.3 Instrument Set-up

12.3.1 Refer to ETS-9-24.0, "Operation and Maintenance of the Micromass Quattro II Triple Quadrupole Mass Spectrometer Fitted with an Atmospheric Pressure Ionization Source," for more details.

12.3.2 Check the solvent level in reservoirs and refill if necessary.

12.3.3 Check the stainless steel capillary at the end of the probe. Use an eyepiece to check the tip. The tip should be flat with no jagged edges. If the tip is found to be unsatisfactory, disassemble the probe and replace the stainless steel capillary.

12.3.4 Turn on the nitrogen.

12.3.5 Open the tune page. Clicks on operate to initiate source block and desolvation heaters.

12.3.6 Open the Inlet Editor.

12.3.6.1 Set HPLC pump to "On"

12.3.6.2 Set the flow to 10 - 500 $\mu\text{L}/\text{min}$ or as appropriate

12.3.6.3 Observe droplets coming out of the tip of the probe. A fine mist should be expelled with no nitrogen leaking around the tip of the probe. Readjust the tip of the probe if no mist is observed

12.3.6.4 Allow to equilibrate for approximately 10 minutes.

12.3.7 The instrument uses these parameters at the following settings. These settings may change in order to optimize the response:

12.3.7.1 Drying gas 250-400 liters/hour

12.3.7.2 ESI nebulizing gas 10-15 liters/hour

12.3.7.3 HPLC constant flow mode flow rate 10 – 500 $\mu\text{L}/\text{min}$

12.3.7.4 Pressure <400 bar (This parameter is not set, it is a guide to ensure the HPLC is operating correctly.)

12.3.7.5 Source block temperature 150°

12.3.7.6 Desolvation temperature 250°

- 12.3.8 Print the tune page, with its parameters, and store it in the study binder with a copy taped into the instrument log.
- 12.3.9 Click on start button in the Acquisition Control Panel (this may vary among MassLynx versions, refer to appropriate MassLynx User's Guide). Ensure start and end sample number includes all samples to be analyzed.

13.0 DATA ANALYSIS AND CALCULATIONS

13.1 Calculations:

13.1.4 Calculate matrix spike percent recoveries using the following equation:

$$\% \text{ Recovery} = \frac{\text{Observed Result} - \text{Background Result}}{\text{Expected Result}} \times 100$$

13.1.5 Calculate percent difference using the following equation:

$$\% \text{ Difference} = \frac{\text{Expected Conc.} - \text{Calculated Conc.}}{\text{Expected Conc.}} \times 100$$

13.1.6 Calculate actual concentrations in matrix (µg/g):

$$\frac{(\text{ng of PFOS calc. from std. Curve} \times \text{Dilution Factor})}{\frac{(\text{Initial Weight of Liver (g)}}{\text{Final Volume (mL)}}} \times \frac{1 \mu\text{g}}{1000 \text{ ng}}$$

14.0 METHOD PERFORMANCE

14.1 Method Detection Limit (MDL) and Limit of Quantitation (LOQ) are method, analyte, and matrix specific. Refer to ETS-8-6.0, Attachment B for a listing of current validated MDL and LOQ values.

14.2 Solvent Blanks, Method Blanks and Matrix Blanks

14.2.1 Solvent blanks, method blanks, and matrix blanks must be below the lowest standard in the calibration curve.

14.3 Calibration Curves

14.3.1 The r^2 value for the calibration must be 0.980 or better.

14.4 Matrix Spikes

14.4.1 Matrix spike percent recoveries must be within $\pm 30\%$ of the spiked concentration.

14.5 Continuing Calibration Verification

14.5.1 Continuing calibration verification percent recoveries must be within $\pm 30\%$ of the spiked concentration.

14.6 If criteria listed in the method performance section are not met, maintenance may be performed on the system and samples reanalyzed or other actions as determined by the analyst. Document all actions in the appropriate logbook.

- 14.7 If data are to be reported when performance criteria have not been met, the data must be footnoted on tables and discussed in the text of the report.

15.0 POLLUTION PREVENTION AND WASTE MANAGEMENT

- 15.1 Sample extract waste and flammable solvent is disposed in high BTU containers, and glass pipette waste is disposed in broken glass containers located in the laboratory.

16.0 RECORDS

- 16.1 Each page generated for a study must have the following information included either in the header or hand written on the page: study or project number, acquisition method, integration method, sample name, extraction date, dilution factor (if applicable), and analyst.
- 16.2 Print the tune page, sample list, and acquisition method from MassLynx to include in the appropriate study folder. Copy these pages and tape into the instrument runlog.
- 16.3 Plot the calibration curve by linear regression, weighted 1/x, then print these graphs and store in the study folder.
- 16.4 Print data integration summary, integration method, and chromatograms from MassLynx and store in the study folder.
- 16.5 Summarize data using suitable software (Excel 5.0+) and store in the study folder, refer to **Attachment A** for an example of a summary spreadsheet.
- 16.6 Back up electronic data to appropriate medium. Record in study notebook the file name and location of backup electronic data.

17.0 TABLES, DIAGRAMS, FLOWCHARTS, AND VALIDATION DATA

- 17.1 Attachment A: ETS-8-7.0 Data summary spreadsheet

18.0 REFERENCES

- 18.1 FACT-M-2.1, "Extraction of Potassium Perfluorooctanesulfonate or Other Fluorochemical Compounds from Liver for Analysis Using HPLC-Electrospray/Mass Spectrometry"
- 18.2 ETS-9-24.0, "Operation and Maintenance of the Micromass Atmospheric Pressure Ionization/Mass Spectrometer Quattro II triple quadrupole Systems"
- 18.3 The validation report associated with this method is ETS-8-6.0 & 7.0-V-1

19.0 AFFECTED DOCUMENTS

- 19.1 ETS-8-6.0, "Extraction of Potassium Perfluorooctanesulfonate or Other Fluorochemical Compounds from Liver or Fluid for Analysis Using HPLC-Electrospray/Mass Spectrometry"

20.0 REVISIONS

Revision
Number

Reason For Revision

Revision
Date

Laboratory Study

Study:

Test Material:

Matrix/Final Solvent:

Method/Revision:

Analytical Equipment System Number:

Instrument Software/Version:

Filename:

R-Squared Value:

Slope:

Y Intercept:

Date of Extraction/Analyst:

Date of Analysis/Analyst:

Group Dose	Sample#	Concentration ng/g	Initial Wt. g	Dilution Factor	Final Conc. ug/g

Slope: Taken from linear regression equation.

Group/Dose: Taken from the study folder.

Sample#: Taken from the study folder.

Concentration (ng/g): Taken from the MassLynx integration summary.

Initial Wt. (g): Taken from the study folder.

Dilution Factor: Taken from the study folder.

Final Conc. (ug/g): Calculated by dividing the initial volume from the concentration

3M ENVIRONMENTAL LABORATORY

METHOD

EXTRACTION OF POTASSIUM PERFLUOROOCTANESULFONATE OR OTHER FLUOROchemical COMPOUNDS FROM LIVER FOR ANALYSIS USING HPLC- ELECTROSPRAY/MASS SPECTROMETRY

Method Number: ETS-8-6.0

Adoption Date:

Revision Date:

Author: Lisa Clemen, Robert Wynne

Approved By:

Laboratory Manager

Date

Group Leader

Date

Technical Reviewer

Date

1.0 SCOPE AND APPLICATION

1.1 Scope: This method is for the extraction of potassium perfluorooctanesulfonate (PFOS) or other fluorochemical compounds from liver.

1.2 Applicable Compounds: Fluorochemical surfactants or other fluorinated compounds.

1.3 Matrices: Rabbit, rat, bovine, and monkey livers or other tissues as designated in the validation report.

2.0 SUMMARY OF METHOD

- 2.1 This method describes the procedure for extracting potassium perfluorooctanesulfonate (PFOS) or other fluorochemical surfactants from liver, or other tissues, using an ion pairing reagent and methyl-*tert*-butyl ether (MtBE). In this method, seven fluorochemicals can be extracted: PFOS, PFOSA, PFOSAA, EtFOSE-OH, PFOSEA, M556, and surrogate standard. An ion pairing reagent is added to the sample and the analyte ion pair is partitioned into MtBE. The MtBE extract is transferred to a centrifuge tube and put onto a nitrogen evaporator until dry. Each extract is reconstituted in 1.0 mL methanol then filtered through a 3 cc plastic syringe attached to a 0.2 μ m nylon filter into glass autovials.
- 2.2 These sample extracts are analyzed following method ETS-8-7.0 or other appropriate methods.

3.0 DEFINITIONS

- 3.1 PFOS: perfluorooctanesulfonate (anion of potassium salt) $C_8F_{17}SO_3$
- 3.2 PFOSA: perfluorooctane sulfonylamide $C_8F_{17}SO_2NH_2$
- 3.3 PFOSAA: perfluorooctane sulfonylamido (ethyl)acetate $C_8F_{17}SO_2N(CH_2CH_3)CH_2CO_2$
- 3.4 EtFOSE-OH: 2(N-ethylperfluorooctane sulfonamido)-ethyl alcohol
 $C_8F_{17}SO_2N(CH_2CH_3)CH_2CH_2OH$
- 3.5 PFOSEA: perfluorooctane sulfonyl ethylamide $C_8F_{17}SO_2N(CH_2CH_3)H$
- 3.6 M556: $C_8F_{17}SO_2N(H)(CH_2COOH)$
- 3.7 Surrogate standard: 1H-1H-2H-2H perfluorooctane sulfonic acid

4.0 WARNINGS AND CAUTIONS

4.1 Health and Safety Warnings:

- 4.1.1 Use universal precautions, especially laboratory coats, goggles, and gloves when handling animal tissue, which may contain pathogens.

5.0 INTERFERENCES

- 5.1 There are no interferences known at this time.

6.0 EQUIPMENT

- 6.1 The following equipment is used while performing this method. Equivalent equipment is acceptable.
- 6.1.1 Ultra-Turrax T25 Grinder for grinding liver samples
- 6.1.2 Vortex mixer, VWR, Vortex Genie 2
- 6.1.3 Centrifuge, Mistral 1000 or IEC
- 6.1.4 Shaker, Eberbach or VWR

6.1.5 Nitrogen Evaporator, Organomation

6.1.6 Balance (sensitivity to 0.100 g)

7.0 SUPPLIES AND MATERIALS

- 7.1 Gloves
- 7.2 Dissecting scalpels
- 7.3 Eppendorf or disposable pipettes
- 7.4 Nalgene bottles, capable of holding 250 mL and 1 L
- 7.5 Volumetric flasks, glass, type A
- 7.6 I-CHEM vials, 40 mL glass
- 7.7 Plastic sample vials, Wheaton, 6 mL (or appropriate size)
- 7.8 Centrifuge tubes, polypropylene, 15 mL
- 7.9 Labels
- 7.10 Oxford Dispensor – 3.0 to 10.0 ml
- 7.11 Syringes, capable of measuring 5 µL to 50 µL
- 7.12 Graduated pipettes
- 7.13 Syringes, disposable plastic, 3 cc
- 7.14 Syringe filters, nylon, 0.2 µm, 25 mm
- 7.15 Timer
- 7.16 Crimp cap autovials and caps
- 7.17 Crimpers

Note: Prior to using glassware and bottles, rinse 3 times with methanol and 3 times with Milli-Q™ water. Rinse syringes a minimum of 9 times with methanol, 3 rinses from 3 separate vials.

8.0 REAGENTS AND STANDARDS

- 8.1 Type I reagent grade water, Milli-Q™ or equivalent; all water used in this method should be Milli-Q™ water and be provided by a Milli-Q TOC Plus™ system
- 8.2 Sodium hydroxide (NaOH), J.T Baker or equivalent
- 8.3 Tetrabutylammonium hydrogen sulfate(TBA), Kodak or equivalent
- 8.4 Sodium carbonate (Na₂CO₃), J.T. Baker or equivalent
- 8.5 Sodium bicarbonate (NaHCO₃), J.T. Baker or equivalent
- 8.6 Methyl-*tert*-butyl ether, Omnisolv, glass distilled or HPLC grade
- 8.7 Methanol, Omnisolv, glass distilled or HPLC grade
- 8.8 Liver, frozen from supplier
- 8.9 Dry ice from supplier
- 8.10 **Fluorochemical standards**
 - 8.10.1 PFOS (3M Specialty Chemical Division), molecular weight = 538

- 8.10.2 PFOSA (3M Specialty Chemical Division), molecular weight = 499
- 8.10.3 PFOSAA (3M Specialty Chemical Division), molecular weight = 585
- 8.10.4 EtFOSE-OH (3M Specialty Chemical Division), molecular weight = 570
- 8.10.5 PFOSEA (3M Specialty Chemical Division), molecular weight = 527
- 8.10.6 M556 (3M Specialty Chemical Division), molecular weight = 557
- 8.10.7 Surrogate standard: 4-H, perfluorooctane sulfonic acid (1-H, 1-H, 2-H, 2-H $C_8F_{13}SO_3H$) molecular weight = 428
- 8.10.8 Other fluorochemicals, as appropriate

8.11 Reagent preparation

NOTE: When preparing larger volumes than listed in reagent, standard, or surrogate preparation, adjust accordingly.

- 8.11.1 10 N sodium hydroxide (NaOH): Weigh approximately 200 g NaOH. Pour into a 1000 mL beaker containing 500 mL Milli-QTM water, mix until all solids are dissolved. Store in a 1 L Nalgene bottle.
- 8.11.2 1 N sodium hydroxide (NaOH): Dilute 10 N NaOH 1:10. Measure 10 mL of 10 N NaOH solution into a 100 mL volumetric flask and dilute to volume using Milli-QTM water. Store in a 125 mL Nalgene bottle.
- 8.11.3 0.5 M tetrabutylammonium hydrogen sulfate (TBA): Weigh approximately 169 g of TBA into a 1 L volumetric containing 500 mL Milli-QTM water. Adjust to pH 10 using approximately 44 to 54 mL of 10 N NaOH (While adding the last mL of NaOH, add slowly because the pH changes abruptly). Dilute to volume with Milli-QTM water. Store in a 1 L Nalgene bottle.
 - 8.11.3.1 TBA requires a check prior to each use to ensure pH = 10. Adjust as needed using 1 N NaOH solution.
- 8.11.4 0.25 M sodium carbonate/sodium bicarbonate buffer ($Na_2CO_3/NaHCO_3$): Weigh approximately 26.5 g of sodium carbonate (Na_2CO_3) and 21.0 g of sodium bicarbonate ($NaHCO_3$) into a 1 L volumetric flask and bring to volume with Milli-QTM water. Store in a 1 L Nalgene bottle.

8.12 Standards preparation

- 8.12.1 Prepare PFOS standards for the standard curve.
- 8.12.2 Prepare other fluorochemical standards, as appropriate. Multicomponent fluorochemical standards are acceptable (for example, one working standard solution containing 1.00 ppm PFOS, 1.02 ppm PFOSA, 0.987 ppm PFOSAA, and 1.10 ppm EtFOSE-OH.)
- 8.12.3 Weigh approximately 100 mg of PFOS into a 100 mL volumetric flask and record the actual weight.
- 8.12.4 Bring to volume with methanol for a stock standard of approximately 1000 ppm ($\mu g/mL$).
- 8.12.5 Dilute the stock solution with methanol for a working standard 1 solution of approximately 50 ppm.

8.12.6 Dilute the stock solution with methanol for a working standard 2 solution of approx. 5.0 ppm.

8.12.7 Dilute the stock solution with methanol for a working standard 3 solution of approx. 0.50 ppm.

8.13 Surrogate stock standard preparation

8.13.1 Weigh approximately 50-60 mg of surrogate standard 1-H, 1-H, 2-H, 2-H, $C_8F_{13}SO_3H$ into a 50 ml volumetric flask and record the actual weight.

8.13.2 Bring to volume with methanol for a surrogate stock of approximately 1000-1200 ppm.

8.13.3 Prepare a surrogate working standard. Transfer approximately 1.0 ml of surrogate stock to a 10 ml volumetric flask and bring to volume with methanol for a working standard of 10-20 ppm. Record the actual volume transferred.

9.0 SAMPLE HANDLING

9.1 All samples are received frozen and must be kept frozen until the extraction is performed.

10.0 QUALITY CONTROL

10.1 Matrix blanks and method blanks

10.1.1 An aliquot of 1.0 mL methanol is used as a solvent blank.

10.1.2 Extract two 1.0 mL aliquots of Milli-Q™ water following this procedure and use as method blanks.

10.1.3 Extract two 1.0 mL aliquots of liver homogenate following this procedure and use as matrix blanks. Refer to 11.1.6.

10.2 Matrix spikes

10.2.1 Prepare and analyze matrix spike and matrix spike duplicate samples to determine the accuracy of the extraction.

10.2.2 Prepare each spike using a sample chosen by the analyst, usually a control liver received with each sample set.

10.2.3 Expected concentrations will fall in the mid-range of the initial calibration curve. Additional spikes may be included and may fall in the low-range of the initial calibration curve.

10.2.4 Prepare one matrix spike and matrix spike duplicate per 40 samples, with a minimum of 2 matrix spikes per batch.

10.3 Continuing calibration verifications

10.3.1 Prepare continuing calibration verification samples to ensure the accuracy of the initial calibration curve.

10.3.2 Prepare, at a minimum, one continuing calibration verification sample per group of 10 samples. For example, if a sample set = 34, four verifications are prepared and extracted.

- 10.3.3 Prepare each continuing calibration verification from the same matrix used to prepare the initial curve.
- 10.3.4 The expected concentrations will fall within the mid-range of the initial calibration curve. Additional spikes may be included that fall in the low-range of the initial calibration curve. This is necessary if the analyst must quantitate using only the low end of the calibration curve (for example, 5 ppb – 100 ppb, rather than 5 ppb – 1000 ppb).

11.0 CALIBRATION AND STANDARDIZATION

11.1 Prepare matrix calibration standards

- 11.1.1 Weigh approximately 40 g of liver into a 250 mL Nalgene bottle containing 200 mLs Milli-Q™ water. Grind to a homogeneous solution.
 - 11.1.2 If 40 g is not available, use appropriate amounts of liver and water to ensure a 1:5 ratio.
 - 11.1.3 Refer to 13.0 to calculate the actual density of liver homogenate and the concentration of solid liver tissue dispersed in 1.0 mL of homogenate solution.
 - 11.1.5 Add 1 mL of homogenate to a 15 mL centrifuge tube. Re-suspend solution by shaking between aliquots while preparing a total of eighteen 1 mL aliquots of homogeneous solution in 15 mL centrifuge tubes.
 - 11.1.6 Two 1 mL aliquots, or other appropriate volume, serve as matrix blanks.
 - 11.1.7 Typically use the standard concentrations and spiking amounts listed in Table 1, at the end of this section, to spike, in duplicate, two standard curves, for a total of eighteen samples, two matrix blanks, and two method blanks.
 - 11.1.8 Refer to validation reports **ETS-8-6.0** and **ETS-8-7.0-V-1** or **Attachment B**, which lists the working ranges and the Linear Calibration Range (LCR) for calibration curves.
 - 11.1.9 Use **Attachment C** as an aid in calculating the concentrations of the working standards. Refer to 13.0 to calculate actual concentrations of PFOS in calibration standards.
- 11.2 To each working standard, blank, or continuing verification, add appropriate amount of surrogate working standard for the concentration to fall within the calibration curve range 5 ppb – 1000ppb.

- 11.3 Extract spiked liver homogenates following 12.14-12.25 of this method. Use these standards to establish each initial curve on the mass spectrometer.

Table 1 Approximate Spiking Amounts for Calibration Standards		
Working Standard (Approx. Conc.)	µl	Approx. final conc. of PFOS in liver
-	-	Blank
0.50 ppm	2	0.005 ppm
0.50 ppm	4	0.010 ppm
0.50 ppm	10	0.025 ppm
0.50 ppm	20	0.050 ppm
0.50 ppm	40	0.100 ppm
5.0 ppm	10	0.250 ppm
5.0 ppm	20	0.500 ppm
5.0 ppm	30	0.750 ppm
50 ppm	4	1.00 ppm

12.0 PROCEDURE

- 12.1 Obtain frozen liver samples.
- 12.2 Cut approximately 1 g of liver using a dissecting scalpel. This part of the procedure is best performed quickly, not allowing the liver to thaw.
- 12.3 Weigh the sample directly into a tared plastic sample vial.
- 12.4 Record the liver weight in the study notebook.
- 12.5 Return unused liver portions to freezer.
- 12.6 Add 2.5 mLs of water to sample vial.
- 12.7 Grind the sample. Put the grinder probe in the sample and grind for about 2 minutes, or until the sample is homogeneous.
- 12.8 Rinse the probe into the sample with 2.5 mLs water using a pipette.
- 12.9 Take the grinder apart and clean it with methanol after each sample. Refer to AMDT-EP-22.
- 12.10 Cap the sample and vortex for 15 seconds. Label the sample vial with the study number, weight, liver ID, date and analyst initials.

- 12.11 Pipette 1.0 mL, or other appropriate volume, of homogenate into a 15 mL polypropylene centrifuge tube. Label the centrifuge tube with the identical information as the sample vial. Refer to attached worksheet for documenting the remaining steps.
- 12.12 Pipette two 1 mL aliquots of Milli-Q™ water to centrifuge tubes. These will serve as method blanks.
- 12.13 Spike all samples, including blanks and standards ready for extraction with surrogate standard as described in section 11.2.
- 12.14 Spike each matrix with the appropriate amount of standard as described in 11.1, or Table 1 of that section, for the calibration curve standards. Also prepare matrix spikes and continuing calibration standards.
- 12.15 Vortex mix the standard curve samples, matrix spike samples, and continuing calibration samples for 15 seconds.
- 12.16 Check to ensure 0.5 M TBA reagent is at pH 10. If not, adjust accordingly.
- 12.17 To each sample, add 1 mL 0.5 M TBA and 2 mL of the 0.25 M sodium carbonate/sodium bicarbonate buffer.
- 12.18 Using an Oxford Dispenser, add 5 mL methyl-*tert*-butyl ether.
- 12.19 Cap each sample and put on the shaker at a setting of 300 rpm, for 20 minutes.
- 12.20 Centrifuge for 20 to 25 minutes at a setting of 3500 rpm, or until layers are well separated.
- 12.21 Label a fresh 15 mL centrifuge tube with the same information as in 12.10.
- 12.22 Remove 4.0 mL of the organic layer to the fresh 15 mL centrifuge tube.
- 12.23 Put each sample on the analytical nitrogen evaporator until dry, approximately 1 to 2 hours.
- 12.24 Add 1.0 mL to each centrifuge tube using a graduated pipette.
- 12.25 Vortex mix for 30 seconds.
- 12.26 Attach a 0.2 µm nylon mesh filter to a 3 cc syringe and transfer the sample to this syringe. Filter into a 1.5 mL glass autovial or low-volume autovial when necessary.
- 12.27 Label the autovial with the study number, animal number and gender, sample timepoint, matrix, final solvent, extraction date, and analyst(s) performing the extraction.
- 12.28 Cap and store extracts at room temperature or at approximately 4 °C until analysis.
- 12.29 Complete the extraction worksheet, attached to this document, and tape in study notebook or include in study binder, as appropriate.

13.0 DATA ANALYSIS AND CALCULATIONS

13.1 Calculations:

- 13.1.1 Calculate the average density of the liver homogenate by recording each mass of ten separate 1.0 mL aliquots of homogenate.

$$\text{Average density (mg/mL)} = \frac{\text{Average mass (mg) of the aliquots}}{1.0 \text{ mL aliquot}}$$

- 13.1.2 Calculate the amount of liver (mg) per 1.0 mL homogenate (or concentration of dispersed solid tissue per mL of homogenate suspension) using the following equation:

$$\frac{\text{g of Liver} \times \text{Average density* of homogenate (mg/mL)}}{(\text{g of Liver} + \text{g of Water})}$$

* refer to 13.1.1 for details.

- 13.1.3 Calculate actual concentrations of PFOS and other fluorochemicals in calibration standards using the following equation:

$$\frac{\mu\text{L of Standard} \times \text{Concentration } (\mu\text{g / mL})}{\text{mg Liver / 1 mL homogenate*}} = \text{Final Concentration } (\mu\text{g/g or mg/kg}) \text{ of PFOS in Liver}$$

*refer to 13.1.2 for details.

14.0 METHOD PERFORMANCE

- 14.1 The method detection limit (MDL) is analyte and matrix specific. Refer to MDL report for specific MDL and limit of quantitation (LOQ) values (refer to **Attachments B and C**).

- 14.2 The following quality control samples are extracted with each batch of samples to evaluate the quality of the extraction and analysis.

14.2.1 Method blanks and matrix blanks.

14.2.2 Matrix spike and matrix spike duplicate samples to determine accuracy and precision of the extraction.

14.2.3 Continuing calibration verification samples to determine the continued accuracy of the initial calibration curve.

- 14.3 Refer to section 14 of ETS-8-7.0 for method performance criteria.

15.0 POLLUTION PREVENTION AND WASTE MANAGEMENT

- 15.1 Sample waste is disposed in biohazard containers, flammable solvent waste is disposed in high BTU containers, and used glass pipette waste is disposed in broken glass containers located in the laboratory.

16.0 RECORDS

- 16.1 Complete the extraction worksheet attached to this method, and tape in the study notebook or include in the 3-ring study binder, as appropriate.

17.0 TABLES, DIAGRAMS, FLOWCHARTS, AND VALIDATION DATA

- 17.1 Attachment A, Extraction worksheet
17.2 Attachment B, MDL/LOQ values and summary
17.3 Attachment C, Calibration standard calculation and concentration worksheet

18.0 REFERENCES

- 18.1 The validation report associated with this method is ETS-8-6.0 & 7.0-V-1.
18.2 AMDT-EP-22, "Routine Maintenance of Ultra-Turrax T-25"
18.3 FACT-M-1.1, "Extraction of PFOS or Other Anionic Fluorochemical Surfactants from Liver for Analysis Using HPLC-Electrospray/Mass Spectrometry"

19.0 AFFECTED DOCUMENTS

- 19.1 ETS-8-7.0, "Analysis of Potassium Perfluorooctanesulfonate or other Fluorochemicals in Liver Extracts using HPLC-Electrospray Mass Spectrometry"

20.0 REVISIONS

<u>Revision Number</u>	<u>Reason For Revision</u>	<u>Revision Date</u>
----------------------------	----------------------------	--------------------------

Cont. Cal. Verifications used the same matrix as for the standard curve.

MDL/LOQ values for rabbit liver

Compound	MDL (ppb)	LOQ (ppb)	Linear Calibration Range (LCR) Approximate concentrations to be used for preparing the Standard Calibration Curve
PFOS	8.45	26.9	30 ppb – 1200 ppb
PFOSA	3.50	11.1	12 ppb – 1200 ppb
PFOSAA	24.6	78.3	30 ppb – 1200 ppb
EtFOSE-OH	108	345	60 ppb – 900 ppb*
M556	82.3	262	60 ppb – 1200 ppb
PFOSEA	33.9	108	30 ppb- 1200 ppb

MDL/LOQ values in rat, bovine, and monkey liver were not statistically determined. Two curves in each of these matrices were extracted and analyzed with the rabbit liver curves to determine equivalence. Responses in the rat, bovine, and monkey liver curves were equivalent to the rabbit responses, therefore, their MDL and LOQ will be assumed to be equivalent to those values as determined for the rabbit liver.

Refer to LOQ Summary and MDL study in ETS-8-6.0 & 7.0-V-1 for further information

* EtFOSE-OH estimates only for MDL and LOQ. Did not meet criteria for validation.

Compound: PFOS

Liver matrix	Prepared range of standards (ppb) (ng/mL)	Range of average curve (ppb) (ng/mL)	LCR from ave curve (ppb) (ng/mL)	Range of low std curve (ppb) (ng/mL)	LCR from low std curve (ppb) (ng/mL)	Range of high std curve (ppb) (ng/mL)	LCR from high std curve (ppb) (ng/mL)
Rabbit	6.19 - 1237	12 - 1200	12 - 1200	6 - 300	12 - 300	60 - 1200	60 - 1200

Compound: PFOSA

Liver matrix	Prepared range of standards (ppb) (ng/mL)	Range of average curve (ppb) (ng/mL)	LCR from ave curve (ppb) (ng/mL)	Range of low std curve (ppb) (ng/mL)	LCR from low std curve (ppb) (ng/mL)	Range of high std curve (ppb) (ng/mL)	LCR from high std curve (ppb) (ng/mL)
Rabbit	6.19 - 1237	12 - 1200	12 - 1200	12 - 300	12 - 300	60 - 1200	60 - 1200

Compound: PFOSAA

Liver matrix	Prepared range of standards (ppb) (ng/mL)	Range of average curve (ppb) (ng/mL)	LCR from ave curve (ppb) (ng/mL)	Range of low std curve (ppb) (ng/mL)	LCR from low std curve (ppb) (ng/mL)	Range of high std curve (ppb) (ng/mL)	LCR from high std curve (ppb) (ng/mL)
Rabbit	6.16 - 1232	12 - 1200	30 - 1200	30 - 900	60 - 900	N/A	N/A

Compound: EtFOSE-OH

Liver matrix	Prepared range of standards (ppb) (ng/mL)	Range of average curve (ppb) (ng/mL)	LCR from ave curve (ppb) (ng/mL)	Range of low std curve (ppb) (ng/mL)	LCR from low std curve (ppb) (ng/mL)	Range of high std curve (ppb) (ng/mL)	LCR from high std curve (ppb) (ng/mL)
Rabbit	6.17 - 1235	31 - 900	31 - 900	N/A	N/A	N/A	N/A

Compound: PFOSEA

Liver matrix	Prepared range of standards (ppb) (ng/mL)	Range of average curve (ppb) (ng/mL)	LCR from ave curve (ppb) (ng/mL)	Range of low std curve (ppb) (ng/mL)	LCR from low std curve (ppb) (ng/mL)	Range of high std curve (ppb) (ng/mL)	LCR from high std curve (ppb) (ng/mL)
Rabbit	6.17 - 1235	31 - 1200	31 - 1200	N/A	N/A	N/A	N/A

Compound: M556

Liver matrix	Prepared range of standards (ppb) (ng/mL)	Range of average curve (ppb) (ng/mL)	LCR from ave curve (ppb) (ng/mL)	Range of low std curve (ppb) (ng/mL)	LCR from low std curve (ppb) (ng/mL)	Range of high std curve (ppb) (ng/mL)	LCR from high std curve (ppb) (ng/mL)
Rabbit	6.17 - 1235	31 - 1200	60 - 1200	N/A	N/A	N/A	N/A

Ion Pair Standard Curves – Tissue

Prep date(s):
 Analyte(s):
 Sample matrix:

Standard number:
 Equipment number:
 Final solvent and TN:
 Blank liver/identifier:

Method/revision:
 Target analyte(s):
 FC mix std approx. 0.500 ppm:
 FC mix std approx. 5.00 ppm:
 FC mix std approx. 50.0 ppm:
 Surrogate std approx. 100 ppm:

Actual concentrations of standards in the FC mix

PFOS Std conc ug/mL	PFOSA Std conc ug/mL	PFOSAA Std conc ug/mL	EtFOSE Std conc ug/mL	PFOSEA Std conc ug/mL	M556 Std conc ug/mL	Std conc ug/mL	All Am't spiked mL	All Density g
0.500	0.500	0.500	0.500	0.500	0.500		0.002	0.167
0.500	0.500	0.500	0.500	0.500	0.500		0.004	0.167
0.500	0.500	0.500	0.500	0.500	0.500		0.010	0.167
0.500	0.500	0.500	0.500	0.500	0.500		0.020	0.167
0.500	0.500	0.500	0.500	0.500	0.500		0.040	0.167
5.00	5.00	5.00	5.00	5.00	5.00		0.010	0.167
5.00	5.00	5.00	5.00	5.00	5.00		0.020	0.167
5.00	5.00	5.00	5.00	5.00	5.00		0.030	0.167
50.0	50.0	50.0	50.0	50.0	50.0		0.004	0.167

Calculated concentrations of standards in the sample matrix

PFOS Final conc ng/g	PFOSA Final conc ng/g	PFOSAA Final conc ng/g	EtFOSE Final conc ng/g	PFOSEA Final conc ng/g	M556 Final conc ng/g	Std conc ng/g	Surrogate Std conc ng/mL	All Am't spiked mL
5.99	5.99	5.99	5.99	5.99	5.99		100	0.005
12.0	12.0	12.0	12.0	12.0	12.0		Surrogate Final conc ng/mL	
29.9	29.9	29.9	29.9	29.9	29.9			
59.9	59.9	59.9	59.9	59.9	59.9			
120	120	120	120	120	120			
299	299	299	299	299	299		0.500	
599	599	599	599	599	599			
898	898	898	898	898	898			
1198	1198	1198	1198	1198	1198			

Validated ranges – approximate concentrations

Liver	PFOS	PFOSA	PFOSAA	EtFOSE-OH	POAA	PFOSEA
Rabbit	5-1000 ppb	5-1000 ppb	5-1000 ppb	5-1000 ppb	5-1000 ppb	5-1000 ppb
Bovine	Estimates only, use rabbit values.					
Rat	Estimates only, use rabbit values.					
Monkey	Estimates only, use rabbit values.					

3M ENVIRONMENTAL LABORATORY

METHOD

ANALYSIS OF POTASSIUM PERFLUOROOCTANESULFONATE OR OTHER FLUOROchemicals IN SERUM EXTRACTS USING HPLC-ELECTROSPRAY/MASS SPECTROMETRY

Method Number: ETS-8-5.1

Adoption Date: 03/01/99

Revision Date:

Author: Lisa Clemen, Robert Wynne

Approved By:

Laboratory Manager

Date

Group Leader

Date

Technical Reviewer

Date

1.0 SCOPE AND APPLICATION

1.1 Scope: This method describes the analysis of serum extracts for fluorochemical surfactants using HPLC-electrospray/mass spectrometry.

1.2 Applicable Compounds: Fluorochemical surfactants or other fluorinated compounds, or other ionizable compounds.

1.3 Matrices: Rabbit, rat, bovine, monkey, and human serum, or other fluids as designated in the validation report.

2.0 SUMMARY OF METHOD

- 2.1** This method describes the analysis of fluorochemical surfactants extracted from serum or other fluids, using HPLC-electrospray/mass spectrometry, or similar system as appropriate. The analysis is performed by monitoring a single ion characteristic of a particular fluorochemical, such as the perfluorooctanesulfonate (PFOS) anion, $m/z = 499$. Additionally, samples may be analyzed using a tandem mass spectrometer to further verify the identity of a compound by detecting daughter ions of the parent ion.

3.0 DEFINITIONS

- 3.1 Atmospheric Pressure Ionization (API):** The Micromass Quattro II triple quadrupole systems allow for various methods of ionization by utilizing various sources, probes, and interfaces. These include but are not limited to: Electrospray Ionization (ESI), Atmospheric Pressure chemical Ionization (APCI), Thermospray, etc. The ionization process in these techniques occurs at atmospheric pressure (i.e., not under a vacuum).
- 3.2 Electrospray Ionization (ES, ESI):** a method of ionization performed at atmospheric pressure, whereby ions in solution are transferred to the gas phase via tiny charged droplets. These charged droplets are produced by the application of a strong electrical field.
- 3.3 Mass Spectrometry, Mass Spectrometer (MS), Tandem Mass Spectrometer (MS/MS):** The API Quattro II triple quadrupole systems are equipped with quadrupole mass selective detectors. Ions are selectively discriminated by mass to charge ratio (m/z) and subsequently detected. A single MS may be employed for ion detection or a series (MS/MS) for more specific fragmentation information.
- 3.4 Conventional vs. Z-spray probe interface:** The latest models of Micromass Quattro II triple quadrupole systems (post 1998) utilize a "Z-spray" conformation. The spray emitted from a probe is orthogonal to the cone aperture. In the conventional conformation it is aimed directly at the cone aperture, after passing through a tortuous pathway in the counter electrode. Though the configuration is different, the methods of operation, cleaning, and maintenance are the same. However, Z-spray components and conventional components are not compatible with one another, but only with similar systems (i.e., Z-spray components are compatible with some other Z-spray systems, etc.)
- 3.5 Mass Lynx Software:** System software designed for the specific operation of these Quattro II triple quadrupole systems. Currently MassLynx has Windows 95 and WindowsNT 4.0 versions. All versions are similar. For more details see the manual specific to the instrument (Micromass Quattro II triple quadrupole MassLynx or MassLynx NT User's Guide).

4.0 WARNINGS AND CAUTIONS

4.1 Health and Safety Warnings:

- 4.1.1** Use caution with the voltage cables for the probe. When engaged, the probe employs a voltage of approximately 5000 Volts.

- 4.1.2 When handling samples or solvents wear appropriate protective gloves, eyewear, and clothing.

4.2 Cautions:

- 4.2.1 Do not operate solvent pumps above capacity of 400 bar (5800 psi) back pressure. If the back pressure exceeds 400 bar, the HP1100 will initiate automatic shutdown.
- 4.2.2 Do not run solvent pumps to dryness.

5.0 INTERFERENCES

- 5.1 To minimize interferences when analyzing samples, teflon should not be used for sample storage or any part of instrumentation that comes in contact with the sample or extract.

6.0 EQUIPMENT

- 6.1 Equipment listed below may be modified in order to optimize the system. Document any modifications in the raw data as method deviations.
 - 6.1.1 Micromass Quattro II triple quadrupole Mass Spectrometer equipped with an electrospray ionization source
 - 6.1.2 HP1100 low pulse solvent pumping system, solvent degasser, column compartment, and autosampler

7.0 SUPPLIES AND MATERIALS

7.1 Supplies

- 7.1.1 High purity grade nitrogen gas regulated to approximately 100 psi (House air system)
- 7.1.2 HPLC analytical column, specifics to be determined by the analyst and documented in the raw data.
- 7.1.3 Capped autovials or capped 15 mL centrifuge tubes

8.0 REAGENTS AND STANDARDS

8.1 Reagents

- 8.1.1 Methanol, HPLC grade or equivalent
- 8.1.2 Milli-Q™ water, all water used in this method should be Milli-Q™ water or equivalent, and may be provided by a Milli-Q TOC Plus system or other vendor
- 8.1.3 Ammonium acetate, reagent grade or equivalent

8.2 Standards

- 8.2.1 Typically two method blanks, two matrix blanks, and eighteen matrix standards are prepared during the extraction procedure. See ETS-8-4.1.

9.0 SAMPLE HANDLING

- 9.1 Fresh matrix standards are prepared with each analysis. Extracted standards and samples are stored in capped autovials or capped 15 mL centrifuge tubes until analysis.
- 9.2 If analysis will be delayed, extracted standards and samples can be refrigerated at approximately 4° C, or at room temperature, until analysis can be performed.

10.0 QUALITY CONTROL

10.1 Solvent Blanks, Method Blanks and Matrix Blanks

10.1.1 Solvent blanks, method blanks and matrix blanks are prepared and analyzed with each batch to determine contamination or carryover.

10.1.2 Analyze a method blank and a matrix blank prior to each calibration curve.

10.2 Matrix Spikes

10.2.1 Matrix spikes are prepared and analyzed to determine the matrix effect on the recovery efficiency.

10.2.2 Matrix spike duplicates are prepared and analyzed to measure the precision and the recovery for each analyte.

10.2.3 Analyze a matrix spike and matrix spike duplicate per forty samples, with a minimum of 2 spikes per batch.

10.2.4 Matrix spike and matrix spike duplicate concentrations will fall in the mid-range of the initial calibration curve. Additional spike concentrations may fall in the low-range of the initial calibration curve.

10.3 Continuing Calibration Verifications

10.3.1 Continuing calibration verifications are analyzed to verify the continued accuracy of the calibration curve.

10.3.2 Analyze a mid-range calibration standard after every tenth sample, with a minimum of one per batch.

11.0 CALIBRATION AND STANDARDIZATION

11.1 Analyze the extracted matrix standards prior to and following each set of extracts. The average of two standard curves will be plotted by linear regression ($y = my + b$), weighted $1/x$, not forced through zero, using MassLynx or other suitable software.

11.2 If the curve does not meet requirements, perform routine maintenance or reextract the standard curve (if necessary) and reanalyze.

11.3 For purposes of accuracy when quantitating low levels of analyte, it may be necessary to use the low end of the calibration curve rather than the full range of the standard curve. Example: when attempting to quantitate approximately 10 ppb of analyte, generate a calibration curve consisting of the standards from 5 ppb to 100 ppb rather than the full range of the curve (5 ppb to 1000 ppb). This will reduce inaccuracy attributed to linear regression weighting of high concentration standards.

12.0 PROCEDURES

12.1 Acquisition Set up

- 12.1.1 Click on start button in the Acquisition Control Panel. Set up a sample list. Assign a filename using MO-DAY-last digit of year-sample number, assign a method (MS) for acquiring, and type in sample descriptions.
- 12.1.2 To create a method click on scan button in the Acquisition control panel and select SIR (Single Ion Recording) or MRM. Set Ionization Mode as appropriate and mass to 499 or other appropriate masses. A full scan is usually collected along with the SIRs. Save acquisition method. If MS/MS instruments are employed, additional product ion fragmentation information may be collected. See Micromass MassLynx GUIDE TO DATA ACQUISITION for additional information and MRM (Multiple Reaction Monitoring).
- 12.1.3 Typically the analytical batch run sequence begins with a set of extracted matrix standards and ends with a set of extracted matrix standards.
- 12.1.4 Samples are analyzed with a continuing calibration check injected after every tenth sample. Solvent blanks should be analyzed periodically to monitor possible analyte carryover and are not considered samples but may be included as such.

12.2 Using the Autosampler

- 12.2.1 Set up sample tray according to the sample list prepared in Section 12.1.1.
- 12.2.2 Set-up the HP1100/autosampler at the following conditions or at conditions the analyst considers appropriate for optimal response. Record actual conditions in the instrument logbook:
- 12.2.2.1 Sample size = 10 μ L injection
- 12.2.2.2 Inject/sample = 1
- 12.2.2.3 Cycle time = 13.5 minutes
- 12.2.2.4 Solvent ramp =

Time	MeOH	2.0 mM Ammonium acetate
0.00 min.	40%	60%
8.50 min.	90%	10%
11.0 min.	90%	10%
12.0 min.	40%	60%

- 12.2.2.5 Press the "Start" button.

12.3 Instrument Set-up

- 12.3.1 Refer to ETS-9-24.0 for more details.
- 12.3.2 Check the solvent level in reservoirs and refill if necessary.

- 12.3.3** Check the stainless steel capillary at the end of the probe. Use an eyepiece to check the tip. The tip should be flat with no jagged edges. If the tip is found to be unsatisfactory, disassemble the probe and replace the stainless steel capillary.
- 12.3.4** Set HPLC pump to "On". Set the flow to 10 - 500 $\mu\text{L}/\text{min}$ or as appropriate. Observe droplets coming out of the tip of the probe. Allow to equilibrate for approximately 10 minutes.
- 12.3.5** Turn on the nitrogen. A fine mist should be expelled with no nitrogen leaking around the tip of the probe. Readjust the tip of the probe if no mist is observed.
- 12.3.6** The instrument uses these parameters at the following settings. These settings may change in order to optimize the response:
- 12.3.6.1** Drying gas 250-400 liters/hour
 - 12.3.6.2** ESI nebulizing gas 10-15 liters/hour
 - 12.3.6.3** HPLC constant flow mode, flow rate 10 - 500 $\mu\text{L}/\text{min}$
 - 12.3.6.4** Pressure <400 bar (This parameter is not set, it is a guide to ensure the HPLC is operating correctly.)
- 12.3.7** Carefully guide the probe into the opening. Insert probe until it will not go any further. Connect the voltage cables to the probe.
- 12.3.8** Print the tune page, with its parameters, and store it in the study binder with a copy taped into the instrument log.
- 12.3.9** Using the cross-flow counter electrode in the ES/MS source is recommended for the analysis of biological matrices.
- 12.3.10** Click on start button in the Acquisition Control Panel (this may vary among MassLynx versions, see appropriate MassLynx USER'S GUIDE). Press the start button. Ensure start and end sample number includes all samples to be analyzed.

13.0 DATA ANALYSIS AND CALCULATIONS

13.1 Calculations:

- 13.1.4** Calculate matrix spike percent recoveries using the following equation:

$$\% \text{ Recovery} = \frac{\text{Observed Result} - \text{Background Result}}{\text{Expected Result}} \times 100$$

- 13.1.5** Calculate percent difference using the following equation:

$$\% \text{ Difference} = \frac{\text{Expected Conc.} - \text{Calculated Conc.}}{\text{Expected Conc.}} \times 100$$

- 13.1.6** Calculate actual concentration of PFOS, or other fluorochemical, in matrix ($\mu\text{g}/\text{mL}$):

$$\frac{(\text{ng of PFOS calc. from std. Curve} \times \text{Dilution Factor})}{(\text{Initial Volume of matrix (mL)} + \text{mL of Surrogate Standard})} \times \frac{1 \mu\text{g}}{1000 \text{ ng}} \times \text{Final Volume (mL)}$$

14.0 METHOD PERFORMANCE

14.1 Method Detection Limit (MDL) and Limit of Quantitation (LOQ) are method, analyte, and matrix specific. Please see ETS-8-4.1, Attachment B, for a listing of current validated MDL and LOQ values.

14.2 Solvent Blanks, Method Blanks, and Matrix Blanks

14.2.1 Solvent blanks, method blanks, and matrix blanks values are must be below the lowest standard in the calibration curve

14.3 Calibration Curves

14.3.1 The r^2 value for the calibration curve must be 0.980 or better.

14.4 Matrix Spikes

14.4.1 Matrix spike percent recoveries are must be within $\pm 30\%$ of the spiked concentration.

14.5 Continuing Calibration Verifications

14.5.1 Continuing calibration verification percent recoveries must be $\pm 30\%$ of the spiked concentration.

14.6 If criteria listed in this method performance section isn't met, maintenance may be performed on the system and samples reanalyzed or other actions as determined by the analyst. Document all actions in the appropriate logbook.

14.7 If data are to be reported when performance criteria have not been met, the data must be footnoted on tables and discussed in the text of the report.

15.0 POLLUTION PREVENTION AND WASTE MANAGEMENT

15.1 Sample extract waste and flammable solvent is disposed in high BTU containers, and glass pipette waste is disposed in broken glass containers located in the laboratory.

16.0 RECORDS

16.1 Each page generated for a study must have the following information included either in the header or hand written on the page: study or project number, acquisition method, integration method, sample name, extraction date, dilution factor (if applicable), and analyst.

16.2 Print the tune page, sample list, and acquisition method from MassLynx to include in the appropriate study folder. Copy these pages and tape into the instrument runlog.

16.3 Plot the calibration curve by linear regression, weighted $1/x$, then print these graphs and store in the study folder.

16.4 Print data integration summary, integration method, and chromatograms, from MassLynx, and store in the study folder.

16.5 Summarize data using suitable software (Excel 5.0) and store in the study folder, see Attachment A for an example of a summary spreadsheet.

- 16.6 Back up electronic data to appropriate medium. Record in study notebook the file name and location of backup electronic data.

17.0 TABLES, DIAGRAMS, FLOWCHARTS, AND VALIDATION DATA

- 17.1 Attachment A: ETS-8-5.1 Data summary spreadsheet.

18.0 REFERENCES

- 18.1 FACT-M-4.1, "Extraction of Potassium Perfluorooctanesulfonate or Other Fluorochemical compounds from Serum for Analysis Using HPLC-Electrospray/Mass Spectrometry"
- 18.2 ETS-9-24.0, "Operation and Maintenance of the Micromass Atmospheric Pressure Ionization/Mass Spectrometer Quattro II triple quadrupole Systems"
- 18.3 The validation report-associated with this method is ETS-8-4.0 & 5.0-V-1.

19.0 AFFECTED DOCUMENTS

- 19.1 ETS-8-4.1, "Extraction of Potassium Perfluorooctanesulfonate or Other Fluorochemical Compounds from Serum for Analysis Using HPLC-Electrospray/Mass Spectrometry"

20.0 REVISIONS

<u>Revision Number</u>	<u>Reason For Revision</u>	<u>Revision Date</u>
1	Section 6.1.2 Clarification of HP1100 system components. Section 11.1 Average of two curves, not standard values, are used for plotting linear regression and added the 1/x weighting of the curve. Section 12.2.2.4 Clarification of solvent ramp. Section 17.1 Changed from attachment B to A.	04/02/99

Laboratory Study

Study:

Test Material:

Matrix/Final Solvent:

Method/Revision:

Analytical Equipment System Number:

Instrument Software/Version:

Filename:

R-Squared Value:

Slope:

Y Intercept:

Date of Extraction/Analyst:

Date of Analysis/Analyst:

Group Dose	Sample#	Concentration ug/mL	Initial Vol. mL	Dilution Factor	Final Conc. ug/mL

Slope: Taken from linear regression equation.

Group/Dose: Taken from the study folder.

Sample#: Taken from the study folder.

Concentration (ug/mL): Taken from the MassLynx integration summary.

Initial Volume (mL): Taken from the study folder.

Dilution Factor: Taken from the study folder.

Final Conc. (ug/mL): Calculated by dividing the initial volume from the concentration

3M ENVIRONMENTAL LABORATORY

METHOD

ANALYSIS OF POTASSIUM PERFLUOROOCTANESULFONATE OR OTHER FLUOROchemicals IN LIVER EXTRACTS USING HPLC-ELECTROSPRAY/MASS SPECTROMETRY

Method Number: ETS-8-7.0

Adoption Date:

Revision Date:

Author: Lisa Clemen, Glenn Langenburg

Approved By:

Laboratory Manager

Date

Group Leader

Date

Technical Reviewer

Date

1.0 SCOPE AND APPLICATION

1.1 Scope: This method is for the analysis of liver extracts for fluorochemical surfactants using HPLC-electrospray/mass spectrometry.

1.2 Applicable Compounds: Fluorochemical surfactants or other fluorinated compounds, or other ionizable compounds.

1.3 Matrices: Rabbit, rat, bovine, monkey liver, or other tissues as designated in the validation report.

2.0 SUMMARY OF METHOD

- 2.1** This method describes the analysis of fluorochemical surfactants extracted from liver using HPLC-electrospray/mass spectrometry, or similar system as appropriate. The analysis is performed by monitoring a single ion characteristic of a particular fluorochemical, such as the perfluorooctanesulfonate (PFOS) anion, $m/z = 499$. Additionally, samples may be analyzed using a tandem mass spectrometer to further verify the identity of a compound by detecting daughter ions of the selected parent ion.

3.0 DEFINITIONS

- 3.1 Atmospheric Pressure Ionization (API):** The Micromass Quattro II triple quadrupole systems allow for various methods of ionization by utilizing various sources, probes, and interfaces. These include but are not limited to: Electrospray Ionization (ESI), Atmospheric Pressure chemical Ionization (APCI), Thermospray, etc. The ionization process in these techniques occurs at atmospheric pressure (i.e. not under a vacuum).
- 3.2 Electrospray Ionization (ES, ESI):** a method of ionization performed at atmospheric pressure, whereby ions in solution are transferred to the gas phase via tiny charged droplets. These charged droplets are produced by the application of a strong electrical field.
- 3.3 Mass Spectrometry, Mass Spectrometer (MS), Tandem Mass Spectrometer (MS/MS):** The API Quattro II triple quadrupole mass spectrometer is equipped with two quadrupole mass selective detectors and a collision cell. Ions are selectively discriminated by mass to charge ratio (m/z) and subsequently detected. A single MS may be employed for ion detection or an ion may be selected in the first quadrupole, fragmented in the collision cell, and these fragments may be analyzed in the second quadrupole.
- 3.4 Conventional vs. Z-spray probe interface:** The latest models of Micromass Quattro II triple quadrupole (post 1998) utilize a "Z-spray" conformation. The spray emitted from a probe is orthogonal to the cone aperture. In the conventional conformation it is aimed directly at the cone aperture, after passing through a tortuous pathway in the counter electrode. Though the configuration is different, the methods of operation, cleaning, and maintenance are the same. However, Z-spray components and conventional components are not compatible with one another, but only with similar systems (i.e. Z-spray components are compatible with other Z-spray systems, etc.)
- 3.5 Mass Lynx Software:** System software designed for the specific operation of these Quattro II triple quadrupole systems. Currently MassLynx has Windows 95 and WindowsNT 4.0 versions. All versions are similar. For more details refer to the manual specific to the instrument (Micromass Quattro II triple quadrupole MassLynx or MassLynx NT User's Guide).

4.0 WARNINGS AND CAUTIONS

4.1 Health and Safety Warnings:

- 4.1.1** Use caution with the voltage cables for the probe. When engaged, the probe employs a voltage of approximately 5000 Volts.

- 4.1.2 When handling samples or solvents wear appropriate protective gloves, eyewear, and clothing.

4.2 Cautions:

- 4.2.1 Operate the solvent pumps below a back pressure of 400 bar (5800 psi). If the back pressure exceeds 400 bar, the HP1100 will initiate automatic shutdown.
- 4.2.2 Do not run solvent pumps to dryness.

5.0 INTERFERENCES

- 5.1 To minimize interferences when analyzing samples, Teflon shall not be used for sample storage or any part of instrumentation that comes in contact with the sample or extract.

6.0 EQUIPMENT

- 6.1 Equipment listed below may be modified in order to optimize the system. Document any modifications in the raw data as method deviations.
 - 6.1.1 Micromass Quattro II triple quadrupole Mass Spectrometer equipped with an electrospray ionization source.
 - 6.1.2 HP1100 low pulse solvent pumping system, solvent degasser, column compartment, and autosampler

7.0 SUPPLIES AND MATERIALS

7.1 Supplies

- 7.1.1 High purity grade air regulated to approximately 100 psi (house air system)
- 7.1.2 HPLC analytical column, specifics to be determined by the analyst and documented in the raw data
- 7.1.3 Capped autovials or capped 15 ml centrifuge tubes

8.0 REAGENTS AND STANDARDS

8.1 Reagents

- 8.1.1 Methanol, HPLC grade or equivalent
- 8.1.2 Milli-Q™ water (ASTM type I), all water used in this method should be ATSM type I, or equivalent, and be provided by a Milli-Q TOC Plus system or other vendor
- 8.1.3 Ammonium acetate, reagent grade or equivalent
 - 8.1.3.1 When preparing different amounts than those listed, adjust accordingly.
 - 8.1.3.2 2.0 mM ammonium acetate solution: Weigh approximately 0.300 g ammonium acetate. Pour into a 2000 mL volumetric container containing 2000 mL Milli-Q™ water, mix until all solids are dissolved. Store at room temperature.

8.2 Standards

- 8.2.1** Typically two method blanks, two matrix blanks, and eighteen matrix standards are prepared during the extraction procedure. Refer to **ETS-8-6.0**.

9.0 SAMPLE HANDLING

- 9.1** Fresh matrix standards are prepared with each analysis. Extracted standards and samples are stored in capped autovials or capped 15 ml centrifuge tubes until analysis.
- 9.2** If analysis will be delayed, extracted standards and samples may be stored at room temperature, or refrigerated at approximately 4° C, until analysis can be performed.

10.0 QUALITY CONTROL

10.1 Method Blanks and Matrix Blanks

- 10.1.1** Solvent blanks, method blanks, and matrix blanks are prepared and analyzed with each batch to determine contamination or carryover.
- 10.1.2** Analyze a method blank and a matrix blank prior to each calibration curve.

10.2 Matrix Spikes

- 10.2.1** Matrix spikes are prepared and analyzed to determine the matrix effect on the recovery efficiency.
- 10.2.2** Matrix spike duplicates are prepared and analyzed to measure the precision and the recovery for each analyte.
- 10.2.3** Analyze a matrix spike and matrix spike duplicate per forty samples. With a minimum of 2 spikes per batch.
- 10.2.4** Matrix spike and matrix spike duplicate concentrations will fall in the mid-range of the initial calibration curve. Additional spike concentrations may fall in the low-range of the initial calibration curve.

10.3 Continuing Calibration Checks

- 10.3.1** Continuing calibration verifications are analyzed to verify the continued accuracy of the calibration curve.
- 10.3.2** Analyze a mid-range calibration standard every tenth sample, with a minimum of one per batch.

11.0 CALIBRATION AND STANDARDIZATION

- 11.1** Analyze the extracted matrix standards prior to and following each set of sample extracts. The average of two standard curves will be plotted by linear regression ($y = mx + b$), weighted $1/x$, not forced through the origin, using MassLynx or other suitable software.
- 11.2** If the curve does not meet requirements perform routine maintenance or reextract the standard curve (if necessary) and reanalyze.

- 11.3** For purposes of accuracy when quantitating low levels of analyte, it may be necessary to use the low end of the calibration curve rather than the full range of the standard curve. Example: when attempting to quantitate approximately 10 ppb of analyte, generate a calibration curve consisting of the standards from 5 ppb to 100 ppb rather than the full range of the curve (5 ppb to 1000 ppb). This will reduce inaccuracy attributed to linear regression weighting of high concentration standards.

12.0 PROCEDURES

12.1 Acquisition Set up

12.1.1 Set up the sample list.

12.1.1.1 Assign a sample list filename using MO-DAY-last digit of year-increasing letter of the alphabet starting with a

12.1.1.2 Assign a method (MS file) for acquiring

12.1.1.3 Assign an HPLC program (Inlet file)

12.1.1.4 Type in sample descriptions and vial position numbers

12.1.2 To create a method click on method in the Acquisition control panel then mass spectrometer headings and select SIR (Single Ion Recording) or MRM (Multiple Reaction Monitoring). Set Ionization Mode as appropriate and mass to 499 or other appropriate masses. A full scan is usually collected along with the SIRs. Save acquisition method. If MS/MS instruments are employed, additional product ion fragmentation information may be collected. Refer to Micromass MassLynx GUIDE TO DATA ACQUISITION for additional information and MRM.

12.1.3 Typically the analytical batch run sequence begins and ends with a set of extracted matrix standards.

12.1.4 Samples are analyzed with a continuing calibration verification injected standard after every tenth sample. Solvent blanks should be analyzed periodically to monitor possible analyte carryover and are not considered samples but may be included as such.

12.2 Using the Autosampler

12.2.1 Set up sample tray according to the sample list prepared in Section 12.1.1.

12.2.2 Set-up the HP1100/autosampler at the following conditions or at conditions the analyst considers appropriate for optimal response. Record actual conditions in the instrument logbook:

12.2.2.1 Sample size = 10 μ L injection

12.2.2.2 Inject/sample = 1

12.2.2.3 Cycle time = 9 minutes

12.2.2.4 Solvent ramp conditions

Time	MeOH	2.0 mM Ammonium acetate
0.00 min.	40%	60%
1.0 min.	40%	60%
4.5 min.	95%	5%
6.5 min.	95%	5%
7.0 min.	40%	60%
9.0 mi.	40%	60%

12.2.2.5 Press the "Start" button.

12.3 Instrument Set-up

12.3.1 Refer to ETS-9-24.0, "Operation and Maintenance of the Micromass Quattro II Triple Quadrupole Mass Spectrometer Fitted with an Atmospheric Pressure Ionization Source," for more details.

12.3.2 Check the solvent level in reservoirs and refill if necessary.

12.3.3 Check the stainless steel capillary at the end of the probe. Use an eyepiece to check the tip. The tip should be flat with no jagged edges. If the tip is found to be unsatisfactory, disassemble the probe and replace the stainless steel capillary.

12.3.4 Turn on the nitrogen.

12.3.5 Open the tune page. Clicks on operate to initiate source block and desolvation heaters.

12.3.6 Open the Inlet Editor.

12.3.6.1 Set HPLC pump to "On"

12.3.6.2 Set the flow to 10 - 500 $\mu\text{L}/\text{min}$ or as appropriate

12.3.6.3 Observe droplets coming out of the tip of the probe. A fine mist should be expelled with no nitrogen leaking around the tip of the probe. Readjust the tip of the probe if no mist is observed

12.3.6.4 Allow to equilibrate for approximately 10 minutes.

12.3.7 The instrument uses these parameters at the following settings. These settings may change in order to optimize the response:

12.3.7.1 Drying gas 250-400 liters/hour

12.3.7.2 ESI nebulizing gas 10-15 liters/hour

12.3.7.3 HPLC constant flow mode flow rate 10 – 500 $\mu\text{L}/\text{min}$

12.3.7.4 Pressure <400 bar (This parameter is not set, it is a guide to ensure the HPLC is operating correctly.)

12.3.7.5 Source block temperature 150°

12.3.7.6 Desolvation temperature 250°

- 12.3.8 Print the tune page, with its parameters, and store it in the study binder with a copy taped into the instrument log.
- 12.3.9 Click on start button in the Acquisition Control Panel (this may vary among MassLynx versions, refer to appropriate MassLynx User's Guide). Ensure start and end sample number includes all samples to be analyzed.

13.0 DATA ANALYSIS AND CALCULATIONS

13.1 Calculations:

13.1.4 Calculate matrix spike percent recoveries using the following equation:

$$\% \text{ Recovery} = \frac{\text{Observed Result} - \text{Background Result}}{\text{Expected Result}} \times 100$$

13.1.5 Calculate percent difference using the following equation:

$$\% \text{ Difference} = \frac{\text{Expected Conc.} - \text{Calculated Conc.}}{\text{Expected Conc.}} \times 100$$

13.1.6 Calculate actual concentrations in matrix ($\mu\text{g/g}$):

$$\frac{(\text{ng of PFOS calc. from std. Curve} \times \text{Dilution Factor})}{\frac{(\text{Initial Weight of Liver (g)})}{\text{Final Volume (mL)}}} \times \frac{1 \mu\text{g}}{1000 \text{ ng}}$$

14.0 METHOD PERFORMANCE

14.1 Method Detection Limit (MDL) and Limit of Quantitation (LOQ) are method, analyte, and matrix specific. Refer to ETS-8-6.0, Attachment B for a listing of current validated MDL and LOQ values.

14.2 Solvent Blanks, Method Blanks and Matrix Blanks

14.2.1 Solvent blanks, method blanks, and matrix blanks must be below the lowest standard in the calibration curve.

14.3 Calibration Curves

14.3.1 The r^2 value for the calibration must be 0.980 or better.

14.4 Matrix Spikes

14.4.1 Matrix spike percent recoveries must be within $\pm 30\%$ of the spiked concentration.

14.5 Continuing Calibration Verification

14.5.1 Continuing calibration verification percent recoveries must be within $\pm 30\%$ of the spiked concentration.

14.6 If criteria listed in the method performance section are not met, maintenance may be performed on the system and samples reanalyzed or other actions as determined by the analyst. Document all actions in the appropriate logbook.

- 14.7 If data are to be reported when performance criteria have not been met, the data must be footnoted on tables and discussed in the text of the report.

15.0 POLLUTION PREVENTION AND WASTE MANAGEMENT

- 15.1 Sample extract waste and flammable solvent is disposed in high BTU containers, and glass pipette waste is disposed in broken glass containers located in the laboratory.

16.0 RECORDS

- 16.1 Each page generated for a study must have the following information included either in the header or hand written on the page: study or project number, acquisition method, integration method, sample name, extraction date, dilution factor (if applicable), and analyst.
- 16.2 Print the tune page, sample list, and acquisition method from MassLynx to include in the appropriate study folder. Copy these pages and tape into the instrument runlog.
- 16.3 Plot the calibration curve by linear regression, weighted 1/x, then print these graphs and store in the study folder.
- 16.4 Print data integration summary, integration method, and chromatograms from MassLynx and store in the study folder.
- 16.5 Summarize data using suitable software (Excel 5.0+) and store in the study folder, refer to **Attachment A** for an example of a summary spreadsheet.
- 16.6 Back up electronic data to appropriate medium. Record in study notebook the file name and location of backup electronic data.

17.0 TABLES, DIAGRAMS, FLOWCHARTS, AND VALIDATION DATA

- 17.1 Attachment A: ETS-8-7.0 Data summary spreadsheet

18.0 REFERENCES

- 18.1 FACT-M-2.1, "Extraction of Potassium Perfluorooctanesulfonate or Other Fluorochemical Compounds from Liver for Analysis Using HPLC-Electrospray/Mass Spectrometry"
- 18.2 ETS-9-24.0, "Operation and Maintenance of the Micromass Atmospheric Pressure Ionization/Mass Spectrometer Quattro II triple quadrupole Systems"
- 18.3 The validation report associated with this method is **ETS-8-6.0 & 7.0-V-1**

19.0 AFFECTED DOCUMENTS

- 19.1 ETS-8-6.0, "Extraction of Potassium Perfluorooctanesulfonate or Other Fluorochemical Compounds from Liver or Fluid for Analysis Using HPLC-Electrospray/Mass Spectrometry"

20.0 REVISIONS

**Revision
Number**

Reason For Revision

**Revision
Date**

Laboratory Study

Study:
 Test Material:
 Matrix/Final Solvent:
 Method/Revision:
 Analytical Equipment System Number:
 Instrument Software/Version:
 Filename:
 R-Squared Value:
 Slope:
 Y Intercept:
 Date of Extraction/Analyst:
 Date of Analysis/Analyst:

Group Dose	Sample#	Concentration ng/g	Initial Wt. g	Dilution Factor	Final Conc. ug/g

Slope: Taken from linear regression equation.
Group/Dose: Taken from the study folder.
Sample#: Taken from the study folder.
Concentration (ng/g): Taken from the MassLynx integration summary.
Initial Wt. (g): Taken from the study folder.
Dilution Factor: Taken from the study folder.
Final Conc. (ug/g): Calculated by dividing the initial volume from the concentration