National Atmospheric Deposition Program

Mercury Analytical Laboratory 2013 Annual Quality Assurance Report

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Table of Contents

1. Q	uality Assurance5
1.1	Philosophy and Objectives5
1.2	Method Detection Limits6
1.3	Accreditations
1.4	Laboratory Bottle Blanks
2.	Quality Control9
2.1	Preparation Blanks
2.2	Initial and Continuing Calibration Verification Standards (ICV & CCVs)12
2.3	Continuing Calibration Blanks14
2.4	Matrix Duplicates
2.5	Matrix Spikes
2.6	Certified Reference Materials
3.	Calculations
3.1	Calculation: Gross MDN Sample Concentration
3.2	Calculation: Net MDN Sample Concentration
3.3	Calculation: MDN Deposition24
4.	Analytical Run Sequence
5.	Proficiency Tests and Laboratory Intercomparison Studies
5.1	Proficiency Tests
6.	Field Quality Control
6.1	Field Bottle Blanks
6.2	Field System Blanks
7.	Quality Rating Codes
8.	Summary and Conclusions
9.	Definitions of Abbreviations and Acronyms
10.	Appendix A: MDL Studies

List of Figures

Figure 1 - 2013 Plot of Total Mercury Mass in Laboratory Bottle Blanks for 105 Samples
Figure 2 - 2013 Plot of Methyl Mercury Mass in Laboratory Bottle Blanks for 12 Samples
Figure 3 - 2013 Control Chart for Total Mercury Concentration in Reagent Preparation Blanks 11
Figure 4 - 2013 Control Chart for Methyl Mercury Concentration in Reagent Preparation Blanks
Figure 5 - 2013 Control Chart for Total Mercury Continuing Calibration Standard Percent Recovery
Figure 6 - 2013 Control Chart for Methyl Mercury Ongoing Calibration Standard Percent Recovery
Figure 7 - 2013 Control Chart for Total Mercury Continuing Calibration Blanks 15
Figure 8 - 2013 Control Chart for Methyl Mercury Continuing Calibration Blanks
Figure 9 - 2013 Control Chart of the Relative Percent Differences for Total Mercury Concentrations in Matrix Duplicates
Figure 10 - 2013 Control Chart of the Relative Percent Differences for Methyl Mercury Concentrations in Matrix Duplicates
Figure 11 - Control Chart for Total Mercury Percent Recovery in Matrix Spikes During 2013 19
Figure 12 - Control Chart for Methyl Mercury Percent Recovery in Matrix Spikes During 2013 . 20
Figure 13 - Control Chart of the Relative Percent Differences for Methyl Mercury Matrix Spike/Matrix Spike Duplicate Pairs during 2013
Figure 14 - Control Chart for Total Mercury Percent Recovery in Certified Reference Material Samples During 2013
Figure 15 - Control Chart for Total Mercury Relative Percent Difference (%RPD) in CRM /CRM Duplicates Samples During 2013
Figure 16 - Control Chart for Methyl Mercury Percent Recovery in Blank Spikes/Blank Spikes Duplicates Samples During 2013
Figure 17 - Control Chart for Methyl Mercury Relative Percent Difference (%RPD) in Blank Spikes/Blank Spikes Duplicates Samples During 2013
Figure 18 - Example of Sample Analysis Worksheet
Figure 19 - Time Series Plot of Total Mercury Concentrations in Field Bottle Blanks During 2013
Figure 20 - Total Mercury Concentration Data for USGS System Blanks and Control Samples During 2013
Figure 21 - Distribution of Quality Rating Codes for Samples Received in 2013

List of Tables

Table 1 - MDL Studies for 2013 Summary Table	7
Table 2 - Laboratory Bottle Blank Summary Table	8
Table 3 - Preparation Blanks Summary Table	. 10
Table 4 - Continuing Calibration Standard Summary Table	. 12
Table 5 - Ongoing Calibration Blanks Summary Table	. 14
Table 6 - Matrix Duplicates Summary Table 2013	. 16
Table 7 - Matrix Spike Recoveries for 2013 Samples	. 19
Table 8 - Matrix Spike/Matrix Spike RPD for 2013 Samples Samples	. 19
Table 9 - Proficiency Tests	. 26
Table 10 - QC Criteria for EPA 1631E and EPA 1630	. 32

Introduction

Eurofins Frontier Global Sciences Inc. (EFGS) has served as the Mercury Analytical Laboratory (HAL) and Site Liaison Center for the Mercury Deposition Network (MDN) since January 1996. MDN, which is coordinated through the National Atmospheric Deposition Program (NADP), was designed with the primary objective of quantifying the wet deposition of mercury in North America to determine long-term geographic and temporal distributions. The MDN consisted of 109 active sites in the United States and Canada at the end of 2013. In 2013, 8 sites were shut down, five new sites were added and two sites were re-started.

The HAL analyzes weekly precipitation samples for total mercury from all active MDN sites and for methyl mercury from 17 sites. The analytical technique, a modified EPA Method 1631, was developed by Nicolas S. Bloom, one of FGS' founders. FGS also served as the referee lab for the EPA Method 1631 "Mercury in Water by Oxidation, Purge and Trap, and Cold Vapor Atomic Fluorescence Spectrometry" final validation study.

EFGS continued to maintain and demonstrate acceptable quality control in 2013. EFGS demonstrated consistency and reproducibility in bottle blanks, preparation blanks, certified reference materials, matrix duplicates, and matrix spikes. Results for all of these quality-control (QC) samples are plotted in control charts and summarized in this report.

The following changes occurred at HAL in 2013:

- Analyst Adela Blaga transferred out of the MDN department in the first quarter. While a new analyst was not hired to replace Adela, some of the remaining analysts were trained on different methods and instruments.
- Analysis of MDN samples for total mercury transferred from the bubbler instruments to the state of the art Tekran 2600 instruments equipped with an autosampler on 9/18/2013.
- Implementation of a new computational model also occurred on 9/18/2013.

1. Quality Assurance

1.1 Philosophy and Objectives

Eurofins Frontier Global Sciences Inc. (EFGS) is committed to a rigorous Quality Assurance program and philosophy. Quality control begins at the bench level. Process improvements are solicited continuously from laboratory technicians and analysts. Management is active in evaluating and implementing feasible improvements. The Quality Assurance program is a system for ensuring that all information, data, and interpretations resulting from an analytical procedure are technically sound, statistically valid, and appropriately documented.

HAL data quality is assessed against EFGS' Data Quality Objectives (DQO). Our DQOs consist of five components: *Precision, Accuracy, Representativeness, Comparability, and Completeness.*

- **Precision** is a measure of data reproducibility. HAL assesses analytical precision using matrix duplicates. The acceptance criterion for matrix duplicates is a relative percent difference (RPD) less or equal to 25 percent (%).
- Accuracy is a measure of proximity to a "true" value. HAL assesses accuracy using certified reference materials and matrix spikes. The acceptance criterion for reference materials and matrix spikes is 75-125% recovery.

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- **Representativeness** is the degree to which a sample's characteristics reflect those of the population. It is demonstrated by accurate, unbiased sampling procedures and appropriate sample processing.
- **Comparability** is measured by comparing the variability of one set of data with respect to another. Control charts enable HAL to assess comparability over the course of an ongoing monitoring project such as MDN.
- **Completeness** is measured by the number of usable data points compared to the number of possible data points. The HAL DQO for the MDN project is at least 95% completeness.

1.2 Method Detection Limits

Method Detection Limits (MDL) are determined according to 40 CFR Part 136, Appendix B. Ten replicates (t-1, 9 degrees of freedom, where t is the Student's T-value for the number of replicates) of matrix-matched samples spiked at 1-10 times the expected MDL are analyzed. There is no recovery criterion for a MDL analysis, but the new calculated MDL value must be within 2 times of the previous established MDL. The standard deviation (σ) is taken from the resulting data and the MDL is determined as t * σ of the replicates. For ten replicates, the MDL is calculated as follows: MDL=2.821 * σ . This value should not be interpreted as the method reporting limit.

The Practical Quantitation Limit (PQL) is the reporting limit for the method and is included as the lowest calibration point (2003 NELAC regulation 5.5.5.2.2.1.h.3 and TNI Standard EL V1M4-2009 section 1.7.1.1.h.iii). The PQL is determined by running ten replicate samples with a concentration that must have the same recovery criteria as for the lowest calibration point.

The ratio between the True Value (TV) and the MDL shall be less than or equal to 10 for a MDL to be valid. A TV/MDL ratio greater than 10 indicates that the study was performed at too high of concentration. In other words, the standard deviation was low at the analyzed level and this does not produce enough variability to establish a realistic MDL. As such, the study would need to be reanalyzed at a lower concentration.

The HAL updates MDL studies periodically for the MDN project. See the summary in Table 1 for the MDL study results performed on the instruments that are used to analyze the MDN samples for total and methyl mercury collected during 2013. All MDL and PQL studies are on file with the Quality Assurance department and are available upon request.

The MDL studies for total mercury for instrument #1 and #9 (datasets THg01-130208-1 and THg09-130208-1), were performed at 0.25 ng/L (the PQL is 0.50 ng/L). The TV/MDL ratios for both instruments were less than 10. Since the TV/MDL ratios were in control for both sets of MDLs, both studies are valid and the higher of the two, 0.118 ng/L, will be used to evaluate data.

Two MDL studies were performed for methyl mercury at a PQL of 0.05 ng/L. A MDL study was performed on instrument #7 in dataset MMHg07-121204-1 and a MDL study was performed on instrument #15 in dataset MMHg15-121204-1. The TV/MDL ratios for both instruments were less than 10. Since the TV/MDL ratios were in control for both sets of MDLs, both studies are valid and the higher of the two, 0.02491 ng/L, will be used to evaluate data.

Instrument	Dataset	MDL (ng/L)	PQL (ng/L)	PQL/MDL
CV-AFS#1	THg01-130208-1	0.034	0.50	14.8
CV-AFS#9	THg09-1302081	0.118	0.50	4.22
CV-GC-AFS #7	MMHg07-121204-1	0.01381	0.050	3.62
CV-GC-AFS #15	MMHg15-121204-1	0.02491	0.050	2.01

Table 1 - MDL Studies for 2013 Summary Table

1.3 Accreditations

Eurofins Frontier Global Sciences currently holds certifications through departments in eight states: California Department of Public Health, Florida Department of Health, State of Louisiana Department of Environmental Quality, State of New York Department of Health, State of New Jersey Department of Environmental Protection, Washington Department of Ecology, Wisconsin Department of Natural Resources, and State of Nevada Division of Natural Resources. Since July 2011, Louisiana's Department of Environmental Quality has been Eurofins Frontier Global Sciences' primary accreditation body for the National Environmental Laboratory Accreditation Program (NELAP). Frontier is also ISO/IEC 17025:2005 and Department of Defense accredited through Perry Johnson Laboratory Accreditation.

1.4 Laboratory Bottle Blanks

1.4.1 Description

Following cleaning, HAL bottles are charged with 20 mL of 1% hydrochloric acid. One sample bottle is randomly selected from each cleaning event and is analyzed for total mercury. On average, 2-3 laboratory bottle blanks are analyzed each week for total mercury. The 20 mL of 1% HCl is oxidized with 1% BrCl, the sample is shaken to ensure that all the walls of the bottles comes in contact with the BrCl. The sample is then left for a minimum of 24 hours before analysis. At least one bottle blank is collected per month and analyzed for methyl mercury.

1.4.2 Purpose

Even in an ultra-clean laboratory, mercury exposure is inherent to the handling of MDN sample bottles. Because such contamination is inevitable, it must be quantified for subtraction from final sample results. Final sample results are corrected by the average bottle blank value from the previous quarter.

1.4.3 Discussion

In 2013, no laboratory bottle blank was higher than the MDL of 0.118 ng/L for total mercury.

In 2013, there were two laboratory bottle blanks that exceed the MDL for methyl mercury (the highest blank at 0.109 ng/bottle doesn't appear on the chart in figure 2). Laboratory bottle blanks are expected to be at, or near, the MDL (0.025 ng/L, Table 2). High bottle blanks are difficult to investigate since there is only enough volume for one analysis and the bottles associated with that batch have already been sent into the field. Possible contamination sources

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are researched, but in these two cases the sources weren't identified. Methyl mercury results are not bottle blank corrected. Note that the values for the bottle blanks are in ng/bottle and the MDL is in ng/L. The bottle blanks are reported as ng/bottle and not ng/L. The laboratory bottle blanks are not converted to ng/L because the 20 mL of the 1% HCl added to the sample bottles is diluted to final volume of water collected at the site. Therefore, the blank values are more meaningful in mass per bottle units.

2013 Laboratory Bottle Blanks	n	Average (ng/bottle)	Standard Deviation	MDL (ng/L)	PQL (ng/L)
Total Mercury	105	0.015	0.014	0.118	0.50
Methyl Mercury	12	0.017	0.031	0.025	0.050

Table 2 - Laboratory Bottle Blank Summary Table

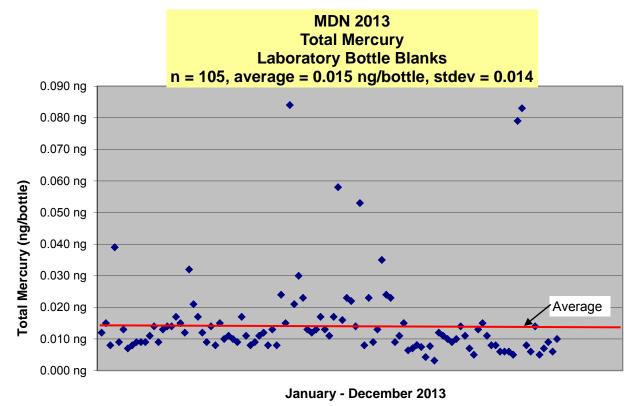


Figure 1 - 2013 Plot of Total Mercury Mass in Laboratory Bottle Blanks for 105 Samples

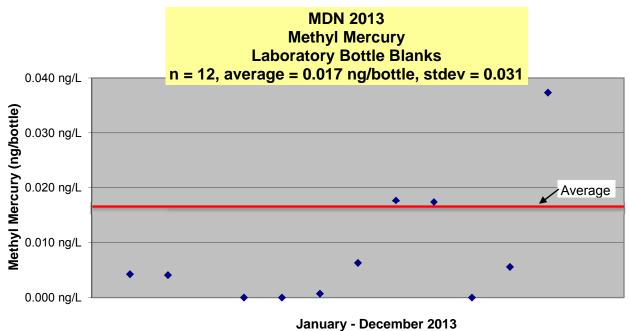


Figure 2 - 2013 Plot of Methyl Mercury Mass in Laboratory Bottle Blanks for 12 Samples

2. Quality Control

QC samples have expected target values that can be used to objectively assess performance of preparation and analytical methods. If performance on these known samples is acceptable, client sample results and other *unknowns* are assumed to be acceptable, as well. Consequently, unacceptable QC results require immediate troubleshooting and re-assessment of affected sample results. The HAL utilizes eight types of QC samples for the MDN project: preparation blanks, continuing calibration standards, continuing calibration blanks, matrix duplicates, matrix spikes, certified reference materials (blank spikes and blank spike duplicates for methyl mercury), field blanks, and system blanks.

2.1 Preparation Blanks

2.1.1 Description

Preparation blanks for total mercury consist of bromine monochloride (1% BrCl), hydroxylamine hydrochloride (0.200 mL), and stannous chloride (0.300 mL) in 100 mL of reagent water. Preparation blanks for methyl mercury consist of 45 mL reagent water, hydrochloric acid (0.4%), ammonium pyrrolidine dithiocarbamate (0.200 mL of APDC) solution, ethylating agent (38.5 μ L) and acetate buffer (0.300 mL). The control limit used at HAL for total mercury is that the absolute value for each individual preparation blank shall be less than 0.25 ng/L. This control limit is lower than the US EPA method 1631E method blank, which individually must be less than 0.50 ng/L, which is the PQL.

The HAL control limit for methyl mercury is set to 0.045 ng/L, which is the same as required by EPA method 1630. See Table 11 for a summary of QC Criteria for EPA 1630 and EPA 1631E.

2.1.2 Purpose

Mercury contamination is inherent in sample preparation and in analytical reagents in any laboratory setting. Preparation blanks are a measure of how much of each sample result can be attributed to these necessary reagents. Preparation blanks also help when investigating possible sources of contamination.

2.1.3 Discussion

In 2013, 4 preparation blanks for total mercury were above the calculated \bar{x} +3 σ limit of 0.128 ng/L. All the preparation blanks analyzed during 2013 were less than the control limit of <0.25 ng/L used at the laboratory and less than the EPA criteria of 0.50 ng/L (table 3 and figure 3).

In 2013, 3 preparation blanks for methyl mercury were above the calculated \bar{x} +3 σ limit of 0.0220 ng/L. None of the preparation blanks were higher than the EFGS control limit of 0.045 ng/L (figure 4). The standard deviation for 2013 of 0.0064 ng/L is less than the EPA requirement of <0.015 ng/L.

High values (greater than \bar{x} +3 σ , but less than control limits) for preparation blanks in August and September don't correspond to a detergent change, equipment change or a newly released set of supplies, but could be caused by samples from an environmental remediation project that was being processed by the lab at that time. Sequestering a set of distillation vials for MDN use only may mitigate this problem. This approach will be attempted in 2015.

2013 Preparation Blanks	n	Average (ng/L)	Stdev (ng/L)	MDL (ng/L)	Mean +3ơ Control Limit (ng/L)	HAL Control Limit (ng/L)	EPA 1631E/1630
Total Mercury	579	0.017	0.037	0.101	0.128	0.25	< 0.50
Methyl Mercury	74	0.0030	0.0064	0.025	0.0220	0.045	Mean <0.045 σ<0.015

Table 3 - Preparation Blanks Summary Table

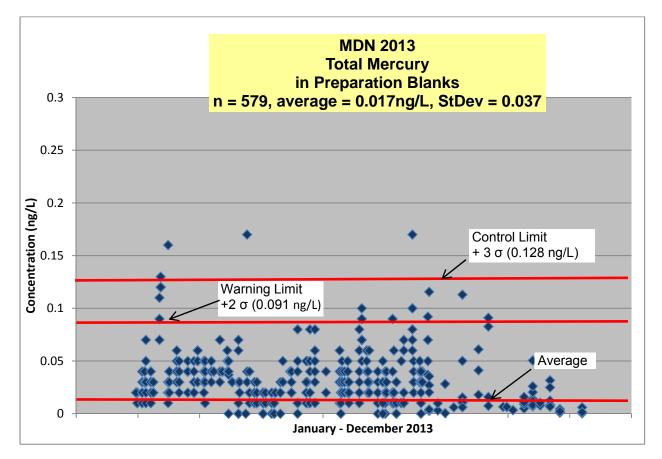


Figure 3 - 2013 Control Chart for Total Mercury Concentration in Reagent Preparation Blanks

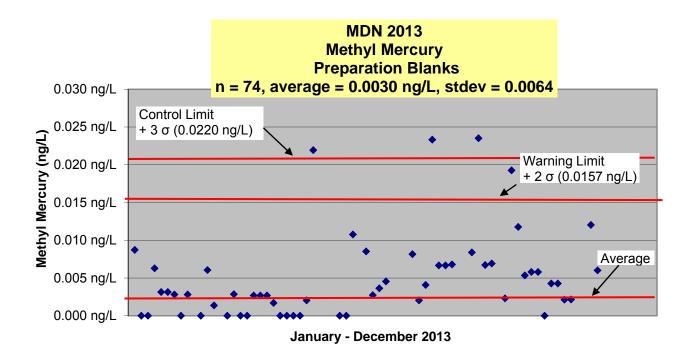


Figure 4 - 2013 Control Chart for Methyl Mercury Concentration in Reagent Preparation Blanks

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Page | 11 Nat

2.2 Initial and Continuing Calibration Verification Standards (ICVs & CCVs)

2.2.1 Description

The Initial Continuing Calibration Verification (ICV) is a solution made from a second source standard, independent of what is used in the primary standard solution. New working standards and standard dilutions are tested prior to use. Three replicates of the new standard are analyzed in the same run as three replicates of the current NIST standard. The mean percent recovery of the three standards should be +/- 5% (95-105%) of the true value and also within 5% of the average NIST recovery. For example, if the average NIST recovery is 97%, the acceptable range for the standards is 95-102%. For the MDN total mercury project, NIST 1641d is the secondary source analyzed after the calibration curve and also after the second set of matrix spikes, and is discussed under the Certified Reference Material (CRM) section.

Continuing Calibration Verification (CCV) standards are analyzed intermittently during the course of sample analysis, after ten or fewer samples, and at the end of each analytical run. The CCV is a standard solution that is made from a traceable stock standard (usually the same source as the primary calibration stock). A 10 ng/L standard for total mercury and a 0.5 ng/L standard for methyl mercury are analyzed as ongoing calibration standards. The MDN control limits for ICVs and CCVs for total mercury are set to 80-120% and for methyl mercury ICVs are 80-120 and CCVs are 75-125%.

2.2.2 Purpose

An ICV is analyzed following each calibration curve to verify the accuracy of the primary standard solution and to validate the calibration curve. CCVs are used to verify that the analytical system is in control or identify analytical drift. All ICV/CCVs reference a unique identification number and are traceable through Frontier's Laboratory Information Management System (LIMS). All raw data reference a unique laboratory ID number and include a unique identifier for each standard used in the analysis.

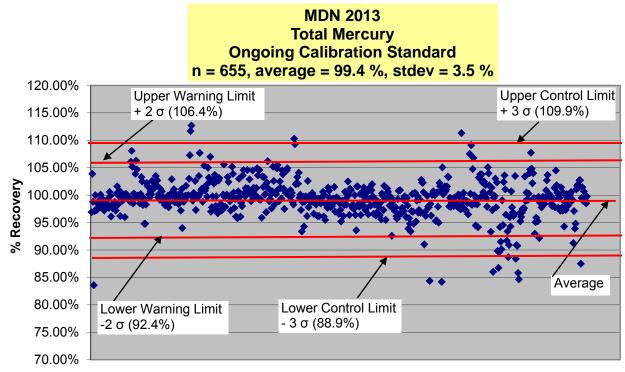
2.2.3 Discussion

Control limits are calculated using the mean value plus/minus three times the standard deviation. For 2013, the range was between 88.9-109.9% for total mercury CCV. Four samples were above the calculated control limit of 109.9% (\bar{x} +3 σ), and eight samples were below the \bar{x} -3 σ of 88.9%. These values were all within the control limit of 80-120% used at HAL (figure 5).

No reportable CCV recoveries were outside the $\bar{x} \pm 3\sigma$ control limit of 51.8-131.7% for methyl mercury (figure 6).

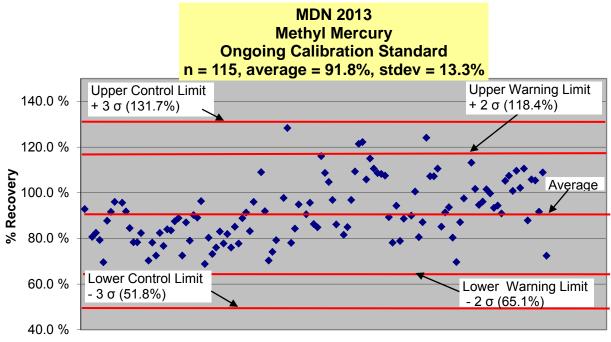
2013 Continuing Calibration Standard	n	Average recovery (%)	Stdev of recovery (%)	±3σ Control Limit (%)	EPA 1631E/1630 Control Limits (%)
Total Mercury	655	99.4	3.5	88.9-109.9	77-123
Methyl Mercury	115	91.8	13.3	51.8-131.7	67-133

Table 4 - Continuing Calibration Standard Summary Table



January - December 2013

Figure 5 - 2013 Control Chart for Total Mercury Continuing Calibration Standard Percent Recovery



January - December 2013

Figure 6 - 2013 Control Chart for Methyl Mercury Ongoing Calibration Standard Percent Recovery

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Page | 13 Nat

National Atmospheric Deposition Program

2.3 Continuing Calibration Blanks

2.3.1 Description

Continuing Calibration Blanks (CCBs) are analyzed during the course of sample analysis, every ten or fewer samples and at the end of each analytical run. Individually, the initial calibration blank (ICB) and each CCB shall be less than 0.25 ng/L in order to be within control limits for total mercury. For MMHg, the mean of the ICB and CCB shall be less than 0.025 ng/L.

2.3.2 Purpose

Instrument blanks are used to monitor baseline drift and to demonstrate freedom from system contamination and carryover.

2.3.3 Discussion

There were several ongoing CCBs for the MDN project in 2013 for total mercury that were outside the calculated control limit of 0.121 ng/L (\bar{x} +3 σ). No CCBs exceeded 0.25 ng/L, which is the control limit that is used for MDN analysis at HAL (table 5 and figure 7).

For 2013, there was one ongoing CCB for methyl mercury that was above the upper control limit of 0.010 ng/L(\bar{x} +3 σ) (table 5 and figure 8). This CCB did not exceed 0.025 ng/L, which is the control limit used for MDN analysis at HAL.

2013 Ongoing Calibration Blanks	n	Average (ng/L)	Stdev (ng/L)	MDL (ng/L)	Upper control Limit +3ơ (ng/L)	EPA 1631E/1630 Control Limits
Total Mercury	1158	0.0043	0.039	0.118	0.121	Individually <0.50 ng/L, mean <0.25 ng/L with a standard deviation <0.10 ng/L
Methyl Mercury	102	0.001	0.003	0.025	0.010	NA

Table 5 - Ongoing Calibration Blanks Summary Table

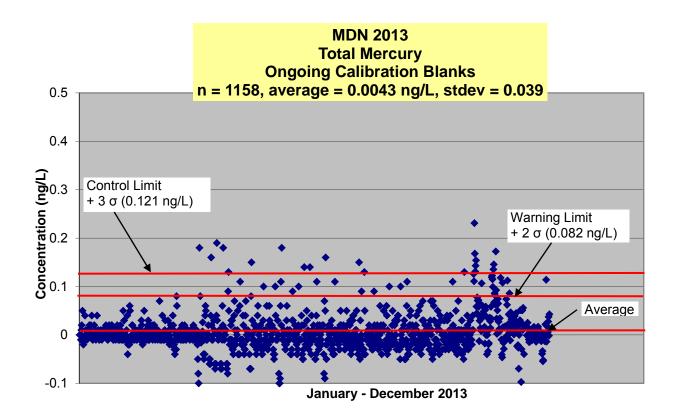


Figure 7 - 2013 Control Chart for Total Mercury Continuing Calibration Blanks

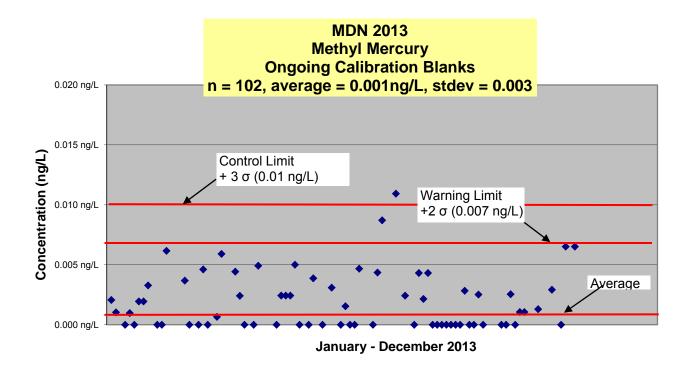


Figure 8 - 2013 Control Chart for Methyl Mercury Continuing Calibration Blanks

2.4 Matrix Duplicates

2.4.1 Description

Matrix Duplicates (MD) are created when an existing sample is split into two portions and then are compared analytically. The MDN control limit for the MDs is set at 25% RPD for total mercury. US EPA methods 1630 and 1631 do not require a MD. One MD is performed for every ten analyzed samples and during a standard MDN THg analytical run, three MDs are analyzed. The source samples are selected depending on available volume. For total mercury analysis, 100 mL is needed for each source sample to obtain the MD, a Matrix Spike (MS), and for potential reanalysis of these QC samples. A smaller aliquot size can be used if needed.

2.4.2 Purpose

Replicate samples provide information about analytical precision. MDs are part of the same sample. As such, their Relative Percent Difference (RPD) is expected to be less than 25%. Out of control results are indications of an inhomogeneous sample matrix and/or poor analytical precision.

2.4.3 Discussion

For 2013, the calculated control limit for total mercury based on \bar{x} +3 σ was 11.9% RPD (table 6). No duplicate samples were above the upper control limit of 25% RPD used at HAL (figure 9).

For methyl mercury, the calculated control limit of $\bar{x} + 3\sigma$ was 40.8% RPD and no duplicate pairs for methyl mercury were above the control limit (table 6 and figure 10). The actual upper control limit used in the laboratory is 25%. For many of the samples, the methyl mercury concentration is lower than, or equal to, the reporting limit of 0.050 ng/L and can result in high RPD.) Also, the recovery criteria for the calibration point at the PQL (0.050 ng/L) level is 70-130%, and analytical values of 0.035 ng/L and 0.065 ng/L, which are within the control criteria for the low calibration point, would be above the acceptance limit of 25% and give a RPD of 60.0%, if these values were produced from duplicate samples. MDN samples of low concentration that produce high RPD values can often be qualified according to the flowcharts used to determine if a qualifier can be applied or not, are included in SOP FGS-038 "Data Review and Validation." HAL applies the same type of qualifiers on MDN data as for any other analysis of EPA 1631 E, if applicable. See Table 12 for qualifiers used at HAL.

Values for QC samples that were qualified for known problems were excluded from the control charts to avoid misrepresentation of actual precision.

2013 Matrix Duplicates	n	Average RPD (%)	Stdev (%)	Upper control Limit +3ơ (%)	EPA 1631E/1630 Control Limits
Total Mercury	605	2.8	3.04	11.9	NA
Methyl Mercury	16	13.5	9.1	40.8	NA

Table 6 - Matrix Duplicates Summary Table 2013

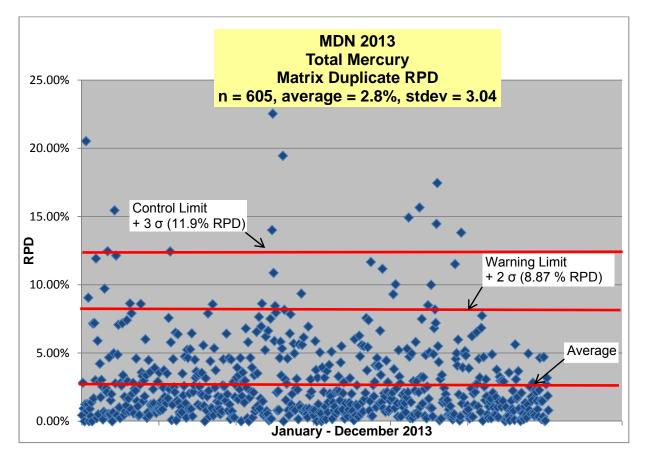


Figure 9 - 2013 Control Chart of the Relative Percent Differences for Total Mercury Concentrations in Matrix Duplicates

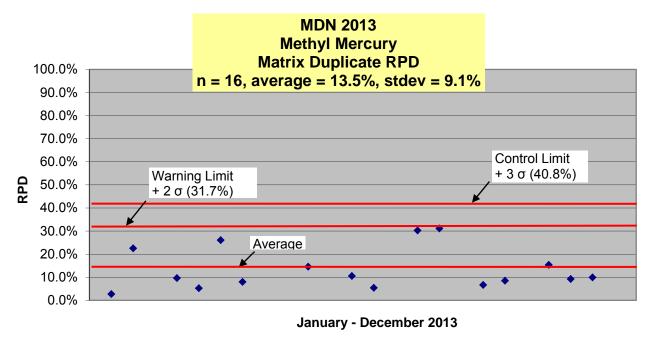


Figure 10 - 2013 Control Chart of the Relative Percent Differences for Methyl Mercury Concentrations in Matrix Duplicates

2.5 Matrix Spikes

2.5.1 Description

A Matrix Spike (MS) for total mercury is created when an MDN sample with known mercury content is split in two fractions and one fraction is supplemented with an additional 1.00 ng of mercury standard.

For both EPA method 1631 and 1630, there must be 1 MS and 1 MSD sample for every 10 samples (a frequency of 10%) and the spiking level shall be at 1–5 times the background concentration or at 1-5 times the MRL (0.5 ng/L for THg and 0.06 ng/L for MMHg), whichever is greater. For MDN runs, due to limited sample volume, only one matrix spike (MS) is performed for every ten analyzed samples and during a normal analytical run three matrix spikes are analyzed. The source samples are selected depending on available volume as 100 mL is desired for the source sample, the matrix duplicate and the matrix spike, and for potential reanalysis of these QC samples. No RPD data for MS/MSD is available for total mercury since only a MS is analyzed. A MS/MSD is performed for methyl mercury and the control limit for the RPD is $\pm 25\%$.

2.5.2 Purpose

The purpose of analyzing a MS and MSD is to demonstrate the performance of the analytical method in a particular sample matrix, and to account for matrix interference. To prepare a MS/MSD, predetermined quantities of the analyte are added to a sample matrix before (when possible) extraction or digestion of samples, in this case preservation with BrCl for total mercury and preservation with HCl and distillation for methyl mercury analysis. If the sample is spiked with the analyte of interest after extraction or digestion, this is considered an analytical spike and an analytical spike duplicate (AS/ASD). If low recovery of a matrix spike is a sign of matrix interference, after investigation by trap and bubbler testing, the samples are diluted and reanalyzed. The purpose is to determine the largest aliquot size that can be analyzed without matrix interference. The source sample is also reanalyzed at the same aliquot size.

2.5.3 Discussion

The control limit for the recovery of the matrix spike for THg based on $\bar{x} \pm 3\sigma$ is 83.4-114.2% (table 7). For 2013, one value was greater than 114.2% and one value was less than 83.4%. All values are within the 75-125% control limit used at HAL (figure 11).

For methyl mercury, a control limit 60.4-143.4% was calculated based on $\bar{x} \pm 3\sigma$ for the recovery of the matrix spike and the matrix spike duplicate (table 7). During 2013, no MS % recovery was greater than the upper control limit of 143.4%. No values were below the lower control limit of 60.4% recovery (figure 12).

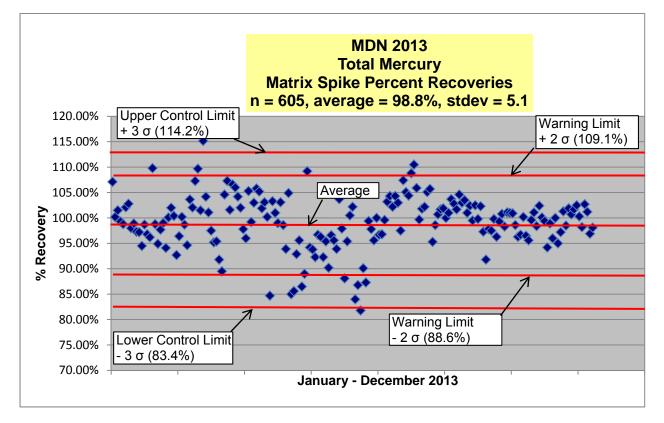
The relative percent difference (RPD) of the methyl mercury matrix spike/matrix spike duplicates had a calculated control limit of 27.7% (\bar{x} +3 σ) for 2013 (table 8). No RPD exceeded this limit, but two RPD values exceeded the 25% control limit used at HAL (figure 13).

2013 Matrix Spikes	n	Average Recovery (%)	Stdev of Recovery (%)	Control Limits ±3σ (%)	HAL Control Limits	EPA 1631E/1630 Control Limits (%)
Total Mercury	605	98.8	5.1	83.4–114.2	75-125	71-125
Methyl Mercury	83	101.9	13.8	60.4-143.4	65-135	65-135

 Table 7 - Matrix Spike Recoveries for 2013 Samples

Table 8 - Matrix Spike/Matrix Spike RPD for 2013 Samples

2013 Matrix Spikes	n	Average RPD (%)	Stdev (%)	+3σ (%)	EPA 1630 Control limits RPD (%)
Methyl Mercury	39	7.6	6.7	27.7	<35%





Eurofins Frontier Global Sciences, Inc. Page | 19

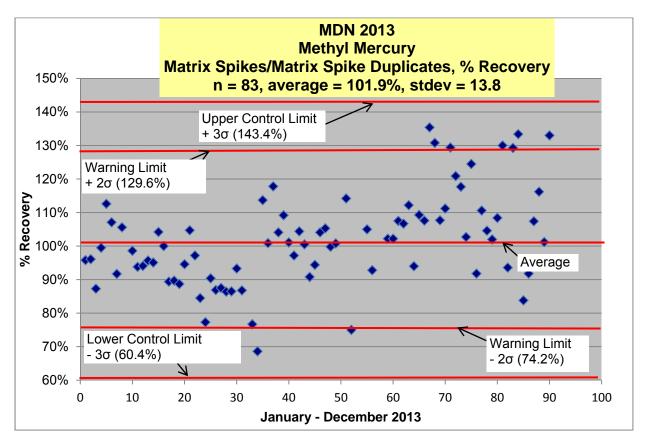


Figure 12 - Control Chart for Methyl Mercury Percent Recovery in Matrix Spikes During 2013

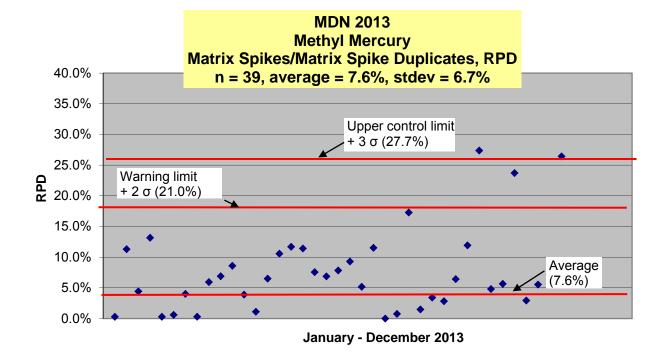


Figure 13 - Control Chart of the Relative Percent Differences for Methyl Mercury Matrix Spike/Matrix Spike Duplicate Pairs during 2013.

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2.6 Certified Reference Materials

2.6.1 Description

Certified Reference Materials (CRMs) are matrix specific standards that are accompanied by a certificate of analysis for the analytes of interest. Eurofins Frontier generally purchases reference materials from the National Institute of Standards and Technology (NIST), the National Research Council of Canada (NRCC), or the International Atomic Energy Agency (IAEA). Eurofins Frontier maintains that matrix equivalent reference materials provide the best measure of precision and accuracy (bias) because they have a consistent, homogeneous matrix.

Currently, there is no available CRM matching the MDN rainwater matrix. Therefore, HAL uses National Institute of Standards and Technology (NIST) reference material 1641d "Mercury in Water." The percent recovery control limits for total mercury are currently set at 75-125% with a RPD of 25%. There is no CRM available for methyl mercury. Therefore, a Blank Spike and a Blank Spike Duplicate (BS/BSD) are analyzed for methyl mercury with acceptance criteria of 70-130%, with a RPD of 25%. The US EPA methods 1630 and 1631 do not require a certified reference material.

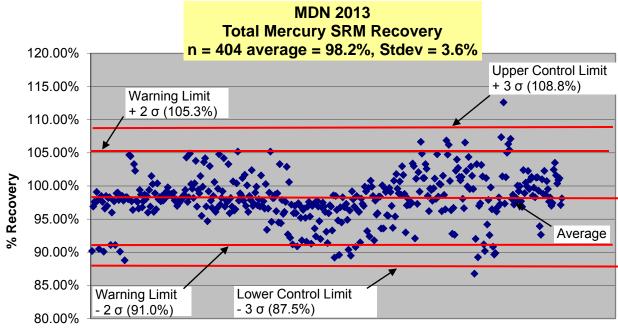
2.6.2 Purpose

Certified Reference Materials are used to demonstrate HAL's ability to recover a target analyte from a specific matrix. For total mercury, the first CRM is analyzed immediately after the calibration standards to validate the analytical curve.

2.6.3 Discussion

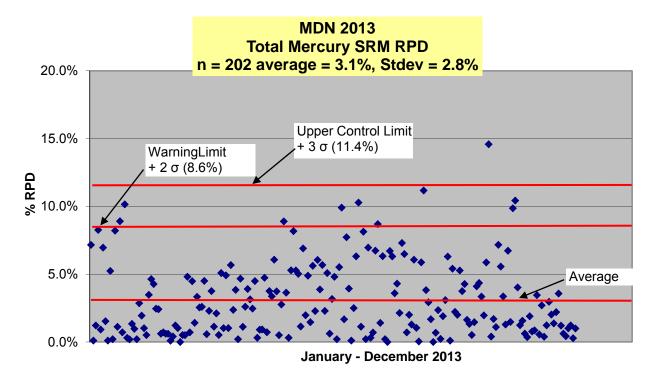
In 2013, the mean of 404 certified reference material recoveries for total mercury was 98.2% with a standard deviation of 3.6% (figure 14). There was one certified reference material recovery above the upper control limit of 108.8% (\bar{x} +3 σ) and one below the lower control limit of 87.5% (\bar{x} -3 σ). All CRM values were within the actual control limit of 75-125% used in the laboratory. The average RPD value for the CRM/CRM duplicate was 3.1% (n=202), with a standard deviation of 2.8%. One RPD values was above the upper control limit calculated by \bar{x} ±3 σ of 11.4%. All of the RPD values were below the 25% limit used in the laboratory, demonstrating good precision between the CRMs and CRM duplicates (figure 15).

In 2013, the mean recovery of 54 blank spikes and blank spike duplicates (BS/BSD) for methyl mercury was 102.8% with a standard deviation of 12.4% (figure 16). No BS % recovery was greater than the upper control limit of 140% based on \bar{x} +3 σ . No values were below the lower control limit of 65.6% recovery based on \bar{x} -3 σ . All values are within the 70-130% control limit used at HAL. The average RPD value for the BS/BSD was 10.3% (n=27) with a standard deviation of 10.0%. All of the RPD values were below the upper control limit calculated by \bar{x} ±3 σ of 39.9%. Three of the RPD values were above the 25% limit used in the laboratory. (figure 17).



January - December 2013







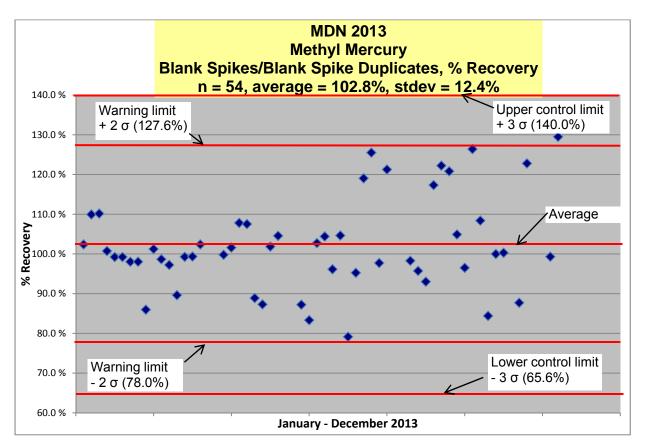


Figure16 - Control Chart for Methyl Mercury Percent Recovery in Blank Spikes/Blank Spikes Duplicates Samples During 2013

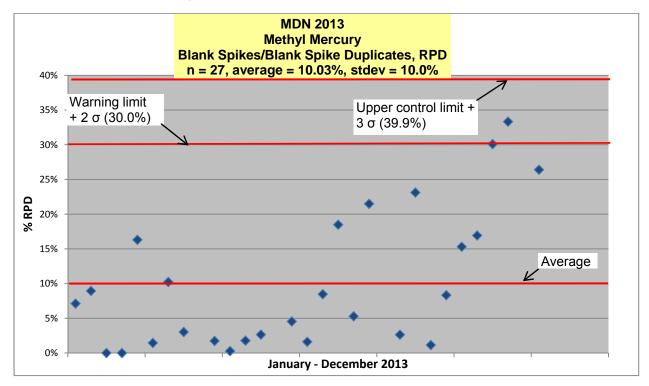


Figure 17 - Control Chart for Methyl Mercury Relative Percent Difference (%RPD) in Blank Spikes/Blank Spikes Duplicates Samples During 2013

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Page | 23

3. Calculations

3.1 Calculation: Gross MDN Sample Concentration

Calc 1){(Sample PA - Ave BB) / Slope} - {(Aliquot * BrCl RB) / 100} = ng Hg/aliquot (mL) Sample PA = sample peak area (PA units) Ave BB = average bubbler blank (PA units) Slope = slope (PA units/ng) Aliquot = volume of sample analyzed (mL) BrCl RB = BrCl reagent blank value (ng/mL of preservative)

1/100 = correction for 1% preservation concentration

3.2 Calculation: Net MDN Sample Concentration

ng Hg/aliquot (mL) * mL / Sample Bottle = ng Hg/Sample Bottle ng Hg/Sample Bottle – ng Hg/Quarterly Bottle Blank = net ng Hg/Sample Bottle net ng Hg/Sample Bottle * (Sample Bottle/mL) * 1000 = net ng Hg/L

3.3 Calculation: MDN Deposition

Deposition = Subppt * Concentration (ng/m^2)

Subppt: Substituted Precip, mm

If on the QA Data Package, "Do Not Use Rain Gage" is not selected, then Subppt is

= RainGauge (inch) * 25.4 (mm/inch)

If this is selected then Subppt is

=BottleCatch (ml) * 25.4 (mm/inch)*0.003281 (inch/mL)

Note: 0.003281 (inch/mL) = comes from 1 inch of capture in sample bottle according to glass funnel opening area of 120 cm² *2.54cm/inch = 304.8 cm³ /inch = 304.8 mL/inch when the density of the rain water is assumed to be 1 g/cm³ = 1 g/mL.

Concentration: Total Hg Concentration in Precipitation

ConcHg = ((sampleHgMass – quarterly BottleBlank) / tmpVol) * 1000

Where:

tmpVol = FullMass – EmptyMass – 20 (20 mL preservative)

SampleHgMass = AliqotHg * (FullMass – EmptyMass) / AliquotVol

4. Analytical Run Sequence

HAL includes the previously mentioned QC samples in all of its analyses for the MDN project. The following work sheet shows how these samples are arranged within a typical analysis day. For every set of ten samples analyzed, the sample set is preceded and followed with a Matrix Duplicate, a Matrix Spike, Continuing Calibration Verification (CCV), and a Continuing Calibration Blank (CCBs). In addition, after the twentieth sample an additional Reference Material sample is analyzed.

	ecipi1 Analysis		n Sample A	nalysis Lab Sheet				FGS D MDN LAB DAT		A SET ID: ET CODE:		
	And	ilyzer: nalyst:		REVIEWER:						DATE:		
alytical Duplicat		sis			S=Samn	e Spike @	Trap Set: 1.00na					
Run	Тр	Bub	HAL Code	Sample ID	ΡΑ	% BrCl	Aliquot Volume	THg per Aliquot	Т	Hg Conc (Net)	Remarks	1
1	1	1		4.00 ng						. ,		۰.
2	2	2		2.00 ng								
3	3	3		1.00 ng								
4	4	4		0.50 ng								
5	5	1		0.05 ng BB-1								•
6 7	6	2 3		BB-2					-			1
8	8	4		BB-3					1			1
9	9	1		NIST1641d		2						1
10	10	2		BrCl-1					\mathbf{T}			1
11	1	3		BrCl-2					T			•
12	2	4		BrCl-3				1	\vdash	Key		
13	3	1		BB-4						- 1		
14	4	2		Sample #1						Defe	wanaa Mata	
15	5	3		Sample #1 D						Refe	rence Mater	Id
16	6	4		Sample #1 S								
17	7	1		Sample #2					1	Drop	aration Blar	100
18	8	2		Sample #3						Prep		IKS
19	9	3		Sample #4								
20	10	4		Sample #5						Matr	ix Duplicate	c
21	1	1		Sample #6						mau	ix Duplicate	Э
22 23	2	2		Sample #7 Sample #8		-						
24	4	4		Sample #9						Matr	ix Spikes	
25	5	1		Sample #10					T	Hau	IX Spines	
26	6	2		1.00								
27	7	3		BB-5						CCV	s	
28	8	4		Sample #11						001	5	
29 30	10	3		Sample #12 Sample #13					+			
31	10	1		Sample #13						CCB	S	
32	2	2		Sample #15						002		-
33	3	3		Sample #16								
34	4	4		Sample #17								1
35	5	1		Sample #18					-			1
36 37	6	2		Sample #19 Sample #20		+			-			1
38	8	4		Sample #11 D					+			1
39	9	3		Sample #11 S					\mathbf{T}			1
40	10	4		1.00								1
41	1	1		BB-6								1
42	2	2		NIST1641d		+			_			4
43 44	3	3		Sample #21 Sample #22		+ +			1			1
44 45	4	4		Sample #22 Sample #23		+		<u> </u>	+			1
46	6	2		etc		1 1		1	1			1
47	7	3							L			1
48	8	4										1
49	9	1				+						1
50	10	2				+			1			1
51 52	1	3			ļ	+			\vdash			1
52	3	4		Sample #21 D		+ +			+			1
54	4	2		Sample #21 S					+			1
55	5	3		1.00				1	\mathbf{t}			1
56	6	4		BB-7				İ	t			1

Figure 18 - Example of Sample Analysis Worksheet

5. Proficiency Tests and Laboratory Intercomparison Studies

Eurofins Frontier Global Sciences participates in two water and two soils pollution proficiency tests each year. One of the water pollution proficiency tests is used for the annual DMR-QA (Discharge Monitoring Report-Quality Assurance) study program, which is a requirement for laboratories that have clients with NPDES (National Pollutant Discharge Elimination System) permits. The Proficiency Test (PT) studies are purchased from a licensed and approved commercial provider. Results for each of these studies are submitted to all of Frontier's accreditation bodies and are available to any client upon request. While these studies are a requirement of accreditation, they are also a valuable tool for internal quality control.

The HAL laboratory is participating in inter-laboratory comparison studies provided by USGS on a monthly basis. Samples are submitted for mercury analysis in both spiked and ultrapure deionized water.

5.1 Proficiency Tests

The proficiency tests listed in table 9 were completed by EFGS during 2013, in addition to the monthly USGS samples that are not included in the table. Results for any tests are available upon request. Control charts for the USGS samples may be viewed at http://bqs.usgs.gov.

Proficiency Test	Organization	Open-close date
ELPAT 85	AIHA	11/4/2013 – 12/3/2013
WP1013	Phenova	10/7/2013 – 11/7/2013
ELPAT 84	AIHA	8/2/2013 – 9/3/2013
IMEP-38	IRMM/JRC/EU	7/3/2013 – 7/30/2013
HW0713	Phenova	7/22/2013 – 9/5/2013
WP0713	Phenova	7/1/2013 – 8/15/2013
ELPAT 83	AIHA	5/1/2013 – 6/3/2013
RR-11522	Phenova	5/17/2013 – 7/1/2013
HW0113	Phenova	1/21/2013 – 3/7/2013
WP0113	Phenova	1/7/2013 – 2/21/2013

Table 9 - Proficiency Tests

6. Field Quality Control

The MDN network utilizes two different procedures to ensure that the sample train is not compromised. The two procedures are field blanks and system blanks.

6.1 Field Bottle Blanks

6.1.1 Description

A field bottle blank has the same contents as a laboratory bottle blank. However, this blank is left exposed at the sampling site for the entire collection period without the collector being opened at any time (no rain accumulation). All field bottle blanks that maintain enough of the initial 20 mL 1% hydrochloric acid (15-21.3 mL) that at least 15 mL can be measured out as aliquot size, are analyzed for total mercury as a field bottle blank sample and are "A" coded and receive "Q" as a sample type. Field blanks with a measured aliquot size less than 15 mL are still analyzed and are "A" coded, but receive "D" (Dry week) as sample type. The analysis is based on mass of sample added to the bubbler and therefore no dilution is needed. There were 55 samples in 2013 that had no recorded precipitation and the event recorder showed the collector did not open, and also had less than 15 mL of preservative in the sample bottle. These results are shown in figure 19. The HAL and the Program Office are attempting to address sample evaporation through lab and field testing. Results from initial testing were submitted at the 2013 NADP Fall Meeting. The HAL will continue its evaporation studies to determine a best practice approach that addresses this issue.

6.1.2 Purpose

Outside of the controlled laboratory environment, the ambient mercury levels increase and this is where the majority of the sample handling occurs. Contamination sources from the surrounding environment are inevitable and their contributions must be quantified so that they can be subtracted from final sample results. High field blanks can be a result of problem with keeping the container closed due to malfunction of the lid seal pad. In dry and windy areas, there is a risk for dust contamination.

6.1.3 Discussion

In 2013, the mean of 121 Field Bottle Blanks was 0.041 ng/bottle with a standard deviation of 0.034 ng/bottle. Figure 19 shows samples NV0220130312, AZ0220130625 and NV0220131112 with elevated mercury values in the field blanks of 0.126, 0.200, 0.291 ng, respectively. NV02 uses an ACM collector, and AZ02 used an NCON Collector. AZ02 had a field blank one month before the high blank that was significantly lower. NV02 had low field blanks within a week of both high blanks. This suggests that neither site had systemic issues of contamination (bad lid seal, handling, etc.). Also, both sites are in the American SW, and are in desert environments. Any windy condition, even if not severe, would have a higher chance of blowing in dust/dirt particles into the sample, which could contribute to a high blank.

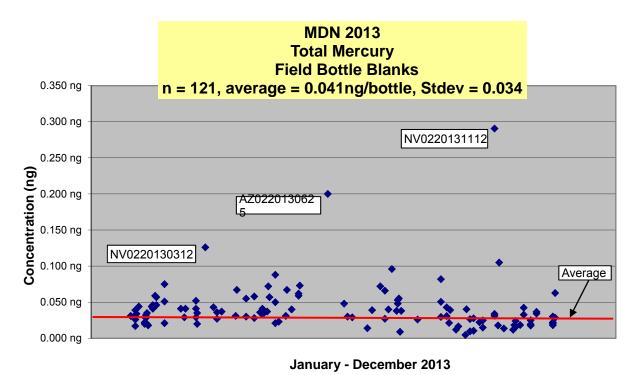


Figure 19 - Time Series Plot of Total Mercury Concentrations in Field Bottle Blanks During 2013

6.2 Field System Blanks

6.2.1 Description

A field system blank is essentially a field bottle blank in which a solution (DI-water) is poured through the wet side collection sample train that was installed in the field for an entire week with no precipitation. The system blank total mercury concentration is compared to the total mercury concentration of an aliquot of the same solution that was not poured through the sample train (i.e. control sample).

6.2.2 Purpose

This quality assurance program, conducted jointly by the U.S. Geological Survey and EFGS, is intended to measure the effects of field exposure, handling, and processing on the chemistry of MDN precipitation samples.

6.2.3 Discussion

In 2013, the mean of 43 system blanks was 0.036 ng/aliquot with a standard deviation of 0.030 ng/aliquot compared to the control sample with a mean of 0.010 ng/aliquot and a standard deviation of 0.019 ng/aliquot. In 2012, the mean of 51 system blanks was 0.073 ng/aliquot with a standard deviation of 0.135 ng/aliquot compared to the control sample with a mean of 0.008 ng/aliquot and a standard deviation of 0.004 ng/aliquot. During 2013, two locations (SC03 and AK06) had higher levels of mercury in their control samples compared to their system blanks. Figure 20 illustrates the system blank results for 2013.

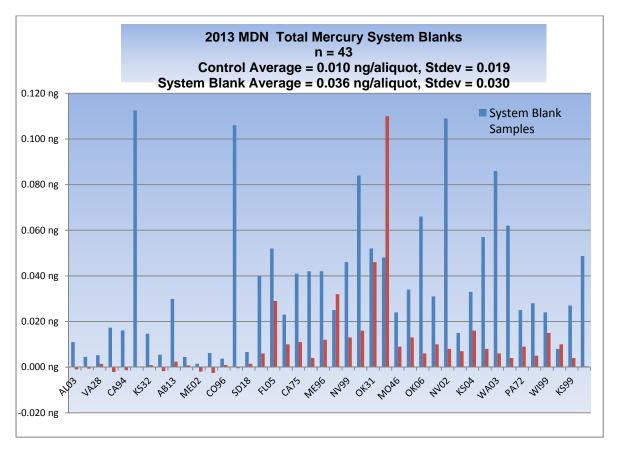


Figure 20 - Total Mercury Concentration Data for USGS System Blanks and Control Samples During 2013

7. Quality Rating Codes

The Quality Rating (QR) code is designed as a user-friendly method to indicate the overall quality of each individual MDN data value. The MDN QR code criterion is modeled after the NADP AIRMON QR code criterion. The QR code is an advisory flag for the general data user. QR codes are assigned by a computer program based on the results of the notes codes given to each MDN sample. Notes codes are defined on the NADP web site at http://nadp.isws.illinois.edu/MDN/mdndata.aspx. A general description of each QR code follows.

A. Valid samples with no problems; contained only precipitation; all sampling and laboratory protocols were followed; all required equipment was installed and operating properly.

B. Valid samples with minor problems; may have contaminants such as insects or other debris; there may be an exception to approved sampling or laboratory methods; required equipment may be lacking or not operating properly. The laboratory does not consider these problems sufficient to invalidate the data, but there is more uncertainty than for A-rated data. These data are used along with A-rated data to calculate average concentrations and deposition.

C. Invalid samples; major problems occurred; the laboratory does not have confidence in the data.

The HAL processed 6008 samples in 2013, which is comparable to the 6234 samples that were processed during 2012. There were 837 samples that received a QR code of "A" while 4739 samples received a "B" QR code, and 432 samples received a "C" QR code. This distribution is illustrated in figure 21. HAL continued to maintain and demonstrate acceptable quality control in 2013. This comparison is based on HAL assessing the QR codes. These codes can later be changed by the NADP Program Office (PO).

Of the 432 "C" coded samples for 2013, 12 incidents occurred at the laboratory.

- 1. AB1420130521: Bubbler blank associated with sample failed high. No volume left for reanalysis
- 2. FL9620130129: Associated matrix duplicate samples failed due to high RPD. No volume left for re-analysis.
- 3. FL97201301229: Associated matrix duplicate samples failed due to high RPD. No volume left for re-analysis.
- 4. KS0420130205: Associated matrix duplicate samples failed due to high RPD. No volume left for re-analysis.
- 5. KS9920130129: Associated matrix duplicate samples failed due to high RPD. No volume left for re-analysis.
- 6. ME0420131204: Sample bottle likely froze and shattered during shipping, and was subsequently transferred to a new bottle. However, transfer wasn't accurately documented and couldn't be completely verified.
- 7. MO0320130122: Associated matrix duplicate samples failed due to high RPD. No volume left for re-analysis.
- 8. OK2220130129: Associated matrix duplicate samples failed due to high RPD. No volume left for re-analysis.
- 9. OK9920130129: Associated matrix duplicate samples failed due to high RPD. No volume left for re-analysis.
- 10. ON0720130521: Bubbler blank associated with sample failed high. No volume left for re-analysis.
- 11. WY2620130122: Associated matrix duplicate samples failed due to high RPD. No volume left for re-analysis.
- 12. WY2620130129: Associated matrix duplicate samples failed due to high RPD. No volume left for re-analysis.

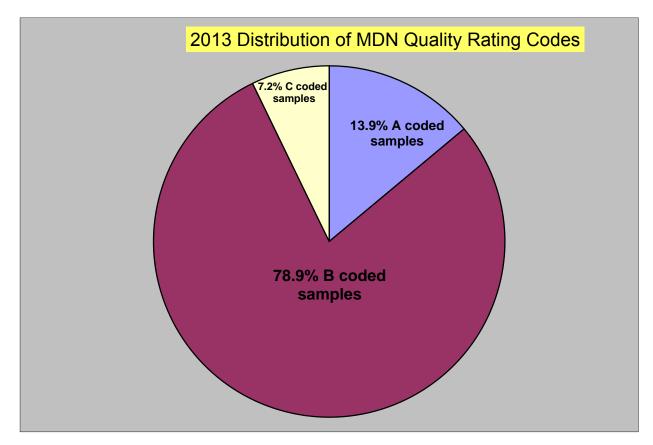


Figure 21 - Distribution of Quality Rating Codes for Samples Received in 2013

8. Summary and Conclusions

The HAL continued to maintain and demonstrate acceptable quality control in 2013. The five DQOs, precision, accuracy, representativeness, comparability, and completeness, were all met. The MDL for total Hg was 0.118 ng/L at a PQL of 0.50 ng/L, and the MDL for MMHg was 0.025 ng/L at a PQL of 0.05 ng/L. Average bottle blank Hg and MMHg content was quantified at 0.015 ng Hg/bottle and 0.017 ng MMHg/bottle, respectively. Preparation and calibration blank total Hg and MHg contents were acceptably low and within control limits. QC sample recoveries for ICVs, CCVs, MS/MSDs, BS/BSDs, and CRMs were within control limits. RPDs for MDs, MSDs, and BSDs were less than $\pm 25\%$. External proficiency testing by Phenova and USGS yielded acceptable results.

Field bottle blanks (n=121) and system blanks (n=43) indicated that field contamination levels continue to be low.

The percentage of C-coded samples increased from 6.6% in 2012 to 7.2% in 2013. The percentage of A-coded samples decreased from 16.4% in 2012 to 13.9% in 2013. The number of B-coded samples increased from 77.0% in 2012 to 78.9% in 2013.

The HAL will continue to look for ways to improve the program both in the laboratory and field to ensure the highest quality data for the MDN.

Table 10 - QC Criteria for EPA 1631E and EPA 1630

QC Item	EPA Method 1631E Criteria	EPA Method 1630 Criteria
Calibration Factor RSD	≤15%	≤15%
Low Standard Recovery	75-125% recovery	65-135% recovery
QCS (Quality Control Sample)	The laboratory must obtain a Quality Control Sample (QCS) from a source different than used to produce the standards. The QCS should be analyzed as an independent check of instrument calibration in the middle of the analytical batch. The recovery criterion is the same as the Ongoing Precision and Recovery (OPR) (77- 123%).	The laboratory must obtain a Quality Control Sample (QCS) from a source different than used to produce the standards. The QCS should be analyzed as an independent check of instrument calibration in the middle of the analytical batch. The recovery criterion is the same as the Ongoing Precision and Recovery (OPR) (77- 123%).
ICV	OPR Standard at 5.0ng/L required at the beginning and end of each run, 77- 123% recovery.	OPR Standard at 0.5ng/L required at the beginning and end of each run, 67-133% recovery.
CCV	No CCV required, see QCS.	No CCV required, see QCS.
MD	No MD required.	No MD required.
MS/MSD	Water: 71-125% Rec. RPD \leq 24% Frequency of 1 MS/MSD per 10 samples. MS/MSD spiking level shall be 1-5 times the sample concentration.	$65-135\%$ recovery with RPD $\leq 35\%$ Frequency of 1 MS/MSD per 10 samples. MS/MSD spiking level shall be 1-5 times the sample concentration.
Bubbler blanks	Individually <0.5ng/L, mean <0.25ng/L with a standard deviation <0.10ng/L. All bubbler blanks are analyzed before the calibration curve.	A single, or more, Ethylation Blanks are analyzed with each analytical run. The value is used to blank correct the standard curve.
ICB and CCB	No ICB, CCBs required.	No ICB, CCBs required.
Preparation Blanks	Minimum of 3, individually <0.50 ng/L.	Minimum of 3. Mean <0.045 ng/L Variability <0.015 ng/L

AIRMoN	Atmospheric Integrated Research Monitoring Network
APDC	Ammonium PyrrolidineDithioCarbamate
AS/ASD	Analytical Spike/ Analytical Spike Duplicate
BS/BSD	Blank Spike/ Blank Spike Duplicate
ССВ	Continued Calibration Blank
CCV	Continued Calibration Verification
CFR	Code of Federal Regulations
CRM	Certified Reference Material
CVAFS	Cold Vapor Atomic Fluorescence Spectrometry
DMR-QA	Discharge Monitoring Report-Quality Assurance
DQO	Data Quality Objectives
EPA	Environmental Protection Agency
EFGS	Eurofins Frontier Global Sciences
HAL	Mercury (Hg) Analytical Laboratory
IAEA	International Atomic Energy Agency
ICB	Initial Calibration Blank
ICV	Initial Calibration Verification
IDL	Instrument Detection Limit
ISO/IEC	International Organization for Standardization (ISO) / International Electrotechnical Commission (IEC)
LCS	Laboratory Control Sample
LCSD	Laboratory Control Sample Duplicate
MD	Matrix Duplicate
MDL	Method Detection Limit
MDN	Mercury Deposition Network
MMHg	Methyl Mercury
MRL	Method Reporting Limit
MS/MSD	Matrix Spike/ Matrix Spike Duplicate
NADP	National Atmospheric Deposition Program
NELAC	National Environmental Laboratory Accreditation Conference
NELAP	National Environmental Laboratory Accreditation Program
NIST	National Institute of Standards and Technology

9. Definitions of Abbreviations and Acronyms

NOS	Network Operations Subcommittee
NPDES	National Pollutant Discharge Elimination System
NRCC	National Research Council Canada
РО	Program Office
PQL	Practical Quantitation Limit
РТ	Proficiency Test
QA	Quality Assurance
QC	Quality Control
QR	Quality Rating
QCS	Quality Control Sample
RPD	Relative Percent Difference
RSD	Relative Standard Deviation
TNI	The NELAC Institute
ТНд	Total Mercury (Hg)
ти	True Value
USGS	United States Geological Survey

10. Appendix A: MDL Studies

Reurofins Frontier Global Sciences	Cold Vapor Atomic Fluorescence Spectroscopy (CV-AFS)
Analyzed by: Jeanne Harrel	Im Anul
Mercury Laboratory Manager: Gerar	rd Van Der Jagt (Jacq 6-11-13
Report Prepared by: Kristine Teffeau	u Ko Teffeau / 5/17/13

Matrix Creation MDL Chudu Tatal Mana

Objective

To establish an Method Detection Limit (MDL) and Practical Quantitation Limit (PQL) for total mercury (THg) in water on CV-AFS 1. This instrument is used to analyze samples received from the Mercury Deposition Network (MDN). Replicates for the MDL study were prepared in accordance with FGS MDN-04.1 and analyzed in accordance with FGS MDN-05.1. The MDL study was performed following the protocols outlined in 40 CFR 136, Appendix B. As detailed below, the MDL for THg in water samples was determined to be **0.034** ng/L THg for CV-AFS 1.

Analytical Method

An instrument calibration was performed according to FGS MDN-05.1. Briefly, this method involves oxidation of the sample through the addition of BrCl, reduction of mercury in the sample aliquot with $SnCl_2$, and analysis by purge and trap and dual amalgamation CV-AFS. The MDL study consisted of the oxidation and analysis of ten water replicates of a 0.250 ng/L solution.

The results for the replicates are listed in the table on the page 2, as well in the raw data sheets for dataset THG01-130208-1. All results are reported <u>corrected</u> for the method and instrument blanks.

MDL Calculation

Using 40 CFR 136, the MDL was calculated using the standard deviation of the spiked replicates. In this case, n = 10 replicates (9 degrees of freedom) and the t value of 2.821 was used at a 99% confidence level. In the following equation, σ is the standard deviation of the results for the spiked replicates at the current PQL.

 $MDL = t^*\sigma = (2.821) * (0.012) = 0.034 \text{ ng/L}.$

MDL and PQL Validation The dataset was peer reviewed and all qualifying parameters (ICV, CCV, CCB, LCS, LCSD, BLKs, R-value, etc.) passed.

A valid MDL requires that the ratio of the spike level (TV) to the calculated MDL must be 10 or less. A valid PQL requires that all replicates must recover within the acceptance criteria specified for the first calibration point (70-130% for this study). The TV/MDL ratio and percent recoveries are listed in the table on page 2. All were in control, making this study valid to establish the MDL and PQL for this analysis and matrix.

\\CUPRUM\General and Admin\Quality Assurance\MDLs & PQLs\MDL Studies (MDN)

MDL Study Data for CV-AFS 1

Dataset ID: THg01-130208-1

Sample	[THg], ng/L		
BrCl-1	0.030		
BrCl-2	0.020		
BrCl-3	0.040		
Mean	0.030		Rec Limits
SD	0.010		70-130%
		Spike Level,	
	Result ng/L	[TV], ng/L	[%Rec]
MDL-Rep1	0.290	0.250	116%
MDL-Rep2	0.290	0.250	116%
MDL-Rep3	0.290	0.250	116%
MDL-Rep4	0.260	0.250	104%
MDL-Rep5	0.270	0.250	108%
MDL-Rep6	0.280	0.250	112%
MDL-Rep7	0.300	0.250	120%
MDL-Rep8	0.270	0.250	108%
MDL-Rep9	0.280	0.250	112%
MDL-Rep10	0.280	0.250	112%
Mean	0.281	0.250	112%
SD	0.012	0.000	4.79%

	[THg], ng/L	Certified Value	[%Rec]
NIST 1641d	7630	7840	97.3%

MDL	0.034
PQL/MDL	7.402

\\CUPRUM\General and Admin\Quality Assurance\MDLs & PQLs\MDL Studies (MDN)

Reurofins Frontier Global Sciences	Matrix Specific MDL Study: Total Mercury in Water by Cold Vapor Atomic Fluorescence Spectroscopy (CV-AFS)		
Analyzed by: <u>Jeanne Harrel</u>	Ann And 66/28/3		
Mercury Laboratory Manager: Gera	ard Van Der Jagt		
Report Prepared by: Kristine Teffeau KETeffeau 6/28/13			

Objective

To establish an Method Detection Limit (MDL) for total mercury (THg) in water on CV-AFS 9. This instrument is used to analyze samples received from the Mercury Deposition Network (MDN). Replicates for the MDL study were prepared in accordance with FGS MDN-04.1 and analyzed in accordance with FGS MDN-05.1. The MDL study was performed following the protocols outlined in 40 CFR 136, Appendix B. As detailed below, the MDL for THg in water samples was determined to be **0.118** ng/L THg for CV-AFS 9.

Analytical Method

An instrument calibration was performed according to FGS MDN-05.1. Briefly, this method involves oxidation of the sample through the addition of BrCl, reduction of mercury in the sample aliquot with SnCl₂, and analysis by purge and trap and dual amalgamation CV-AFS. The MDL study consisted of the oxidation and analysis of ten water replicates of a 0.250 ng/L solution.

The results for the replicates are listed in the table on the page 2, as well in the raw data sheets for dataset THG09-130208-1. All results are reported <u>corrected</u> for the method and instrument blanks.

MDL Calculation

Using 40 CFR 136, the MDL was calculated using the standard deviation of the spiked replicates. In this case, n = 10 replicates (9 degrees of freedom) and the t value of 2.821 was used at a 99% confidence level. In the following equation, σ is the standard deviation of the results for the spiked replicates at the current PQL.

 $MDL = t^*\sigma = (2.821) * (0.042) = 0.118 \text{ ng/L}.$

MDL and PQL Validation The dataset was peer reviewed and all qualifying parameters (ICV, CCV, CCB, LCS, LCSD, BLKs, R-value, etc.) passed.

A valid MDL requires that the ratio of the spike level (TV) to the calculated MDL must be 10 or less. A valid PQL requires that all replicates must recover within the acceptance criteria specified for the first calibration point (70-130% for this study). The TV/MDL ratio and percent recoveries are listed in the table on page 2. The ratio was in control, making this study valid to establish the MDL for this analysis and matrix. One of the percent recoveries did not fall within limits. This study does not verify the PQL of 0.250 ng/L.

MDL Study Data for CV-AFS 9

Dataset ID: THg09-130208-1

Sample	[THg], ng/L		
BrCl-1	0.030		
BrCl-2	0.020		
BrCl-3	0.040		
Mean	0.030		Rec Limits
SD	0.010		70-130%
		Spike Level,	
	Result ng/L	[TV], ng/L	[%Rec]
MDL-Rep1	0.310	0.250	124%
MDL-Rep2	0.310	0.250	124%
MDL-Rep3	0.270	0.250	108%
MDL-Rep4	0.250	0.250	100%
MDL-Rep5	0.170	0.250	68.0%
MDL-Rep6	0.240	0.250	96.0%
MDL-Rep7	0.300	0.250	120%
MDL-Rep8	0.250	0.250	100%
MDL-Rep9	0.280	0.250	112%
MDL-Rep10	0.280	0.250	112%
Mean	0.266	0.250	106%
SD	0.042	0.000	16.78%
	[THg], ng/L	Certified Value	[%Rec]
NIST 1641d	7120	7840	90.8%

MDL	0.118	
PQL/MDL	2.112	

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Analyzed by: Adela Blaga	DIHG 01117/13
Mercury Supervisor: Ryan Nelson	Kn MM 1/19/13
Report Prepared by: Ryan Nelson	Mr. N/ 1/13/13
Report Reviewed by: Kristine Teffeau	6 Tekleau 1/16/13

MDL Study Data for Methyl Mercury in Precipitation Samples Preparation Method: FGS-MDN-07 Analysis Method: FGS-MDN-08 Dataset: MHG7-121204-1

Objective To verify the existing Method Detection Limit (MDL). This MDL verification study applies to Methyl Mercury in precipitation samples as prepared by method EFGS-013 and analyzed by method EFGS-070. The MDL verification was calculated and evaluated in accordance with 40 CFR 136. As detailed below, the MDL for Methyl Mercury in Water was determined to be **0.01381 ng/L MHg.**

<u>Analytical Method</u> Before analysis, the MHg in an aliquot of sample is co-distilled into pure water. The distillates are then analyzed for MHg by aqueous phase ethylation and Cold Vapor – Gas Chromatography – Atomic Fluorescence Spectroscopy (CV-GC-AFS).

For this dataset, an efficiency factor of 0.8046 was used and the calibration was performed according to FGS-MDN-08.

The MDL study consisted of a 0.050 ng/L solution of MHg divided into ten replicates, which were distilled then analyzed. The results of these measurements are found in the table on the next page, as well in the raw data sheets. All peak heights were **corrected** for the instrument blanks (0.30 units). All final concentrations were **corrected** for the preparation blanks (0.000 units).

<u>MDL</u> Calculation Using 40 CFR 136, the MDL was calculated using the standard deviation of the spiked samples, with n = 10 replicates (9 degrees of freedom). In this case, the t value of 2.821 was used in the following equation, where σ is the standard deviation of the results obtained on the replicates.

$MDL = t^*\sigma$

The MDL calculated from these data is (2.821)*(0.00438), or **0.01381 ng/L MHg**.

MDL Validation The dataset was peer reviewed and all qualifying parameters (ICV, CCV, CCB, LCS, R-value, etc.) passed. An existing MDL is verified if a newly-calculated MDL is less than 2x the existing value and the new study yields a PQL/MDL ratio which does not exceed 10. The existing Methyl Mercury MDL of 0.026 ng/L was verified and the MDL did not require updating in the LIMS. See table below.

Percent recoveries for the studies are evaluated only when a new PQL needs to be established. The criteria for the percent recoveries are the acceptance limits for the low calibration point (70-130%). All 10 replicates were spiked at the current PQL (0.050 ng/L) and recovered between 70-130%.

12/4/2012 MMHg07-121204-1 MDN MDL Study

Eurofins Frontier Global Sciences 11720 North Creek Pkwy North, Suite 400 Bothell, WA 98011

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Sample	[MHg], ng/L	6	
F212003-BLK1	0.000		
F212003-BLK2	-0.005		
F212003-BLK3	-0.005		
Mean	-0.003		
SD	0.003		70-130%
	[MHg], ng/L	[TV], ng/L	[%Rec]
F212003-BS2	0.053	0.050	106%
F212003-BS3	0.053	0.050	106%
F212003-BS4	0.046	0.050	92.0%
F212003-BS5	0.043	0.050	86.0%
F212003-BS6	0.058	0.050	116%
F212003-BS7	0.046	0.050	92.0%
F212003-BS8	0.048	0.050	96.0%
F212003-BS9	0.043	0.050	86.0%
F212003-BSA	0.046	0.050	92.0%
F212003-BSB	0.046	0.050	92.0%
Mean	0.048	0.050	96.4%
SD	0.005	0.000	9.8%

F202026-BS1	0.90	1.00	89.9%	%RPD
F202026-BSD1	1.02	1.00	102%	12.9%

MDL	0.01381
TV/MDL	3.621
FMDL in LIMS	0.026
2x LIMS FMDL	0.052
FPQL in LIMS	0.050

\Quality Assurance\MDLs & PQLs\MDL Study (MHg)\MDL Verification MHg Water CV-GC-AFS #7 12.4.12

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Analyzed by: Adela Blaga / 90/	01/17/13
Mercury Supervisor: Ryan Nelson	1 N 4 1/17/13
Report Prepared by: Ryan Nelson	12 Nr 1/17/13
Report Reviewed by: Kristine Teffeau	KE Teffeau 1/17/13

MDL Study Data for Methyl Mercury in Precipitation Samples Preparation Method: FGS-MDN-07 Analysis Method: FGS-MDN-08 Dataset: MHG15-121204-1

Objective To verify the existing Method Detection Limit (MDL). This MDL verification study applies to Methyl Mercury in precipitation samples as prepared by method EFGS-013 and analyzed by method EFGS-070. The MDL verification was calculated and evaluated in accordance with 40 CFR 136. As detailed below, the MDL for Methyl Mercury in Water was determined to be **0.02491 ng/L MHg.**

<u>Analytical Method</u> Before analysis, the MHg in an aliquot of sample is co-distilled into pure water. The distillates are then analyzed for MHg by aqueous phase ethylation and Cold Vapor – Gas Chromatography – Atomic Fluorescence Spectroscopy (CV-GC-AFS).

For this dataset, an efficiency factor of 0.8046 was used and the calibration was performed according to FGS-MDN-08.

The MDL study consisted of a 0.050 ng/L solution of MHg divided into ten replicates, which were distilled then analyzed. The results of these measurements are found in the table on the next page, as well in the raw data sheets. All peak heights were **corrected** for the instrument blanks (0.10 units). All final concentrations were **corrected** for the preparation blanks (0.005 units).

MDL Calculation Using 40 CFR 136, the MDL was calculated using the standard deviation of the spiked samples, with n = 10 replicates (9 degrees of freedom). In this case, the t value of 2.821 was used in the following equation, where σ is the standard deviation of the results obtained on the replicates.

$MDL = t^*\sigma$

The MDL calculated from these data is (2.821)*(0.00438), or **0.02491 ng/L MHq**.

MDL Validation The dataset was peer reviewed and all qualifying parameters (ICV, CCV, CCB, LCS, R-value, etc.) passed. An existing MDL is verified if a newly-calculated MDL is less than 2x the existing value and the new study yields a PQL/MDL ratio which does not exceed 10. The existing Methyl Mercury MDL of 0.026 ng/L was verified and the MDL did not require updating in the LIMS. See table below.

Percent recoveries for the studies are evaluated only when a new PQL needs to be established. The criteria for the percent recoveries are the acceptance limits for the low calibration point (70-130%). All 10 replicates were spiked at the current PQL (0.050 ng/L) and recovered between 70-130%.

12/4/2012 MMHg15-121204-1 MDN MDL Study

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Sample	[MHg], ng/L		
F212006-BLK1	0.013		
F212006-BLK2	0.003		
F212006-BLK3	0.003		
Mean	0.006		
SD	0.006		70-130%
	[MHg], ng/L	[TV], ng/L	[%Rec]
F212006-BS2	0.035	0.050	70.0%
F212006-BS3	0.035	0.050	70.0%
F212006-BS4	0.038	0.050	76.0%
F212006-BS5	0.056	0.050	112%
F212006-BS6	0.035	0.050	70.0%
F212006-BS7	0.056	0.050	112%
F212006-BS8	0.038	0.050	76.0%
F212006-BS9	0.041	0.050	82.0%
F212006-BSA	0.035	0.050	70.0%
F212006-BSB	0.051	0.050	102%
Mean	0.042	0.050	84.0%
SD	0.009	0.000	17.7%

F202026-BS1	1.00	1.00	100%	%RPD
F202026-BSD1	1.17	1.00	117%	15.7%

MDL	0.02491
TV/MDL	2.007
FMDL in LIMS	0.026
2x LIMS FMDL	0.052
FPQL in LIMS	0.050

\Quality Assurance\MDLs & PQLs\MDL Study (MHg)\MDL Verification MHg Water CV-GC-AFS #15 12.4.12