

Biological Analysis Report

NE-DIST-2023-11-16-01

Florida DEP Laboratory
2600 Blair Stone Road
Tallahassee, FL 32399-2400
DOH Accreditation E31780

Overflow Analyses Performed By:
Diversified Environmental Laboratories, Inc.
3653 Regent Boulevard
Suite 509
Jacksonville, FL 32224
DOH Accreditation E821059

Event Description: **Ormond Boat and L.Eaton**
Request ID: **RQ-2023-11-27-24**
Customer: **NE-DIST**
Project ID: **SMP**

Send Reports to:
FL Dept. of Environmental Protection
8800 Baymeadows Way West
Suite 100
Jacksonville, FL 32256
Attn: Jeremy Parrish

For additional information please contact
Cheryl Swanson - Administrator
Sarah Menz, Ph.D. - Bench Biology & Aquatic
Toxicity
Puja Jasrotia, Ph.D. - Molecular Biology & Taxonomy
Thekkekalathil Chandrasekhar, Ph.D, QA Officer
Phone (850) 245-8177

Certified by: Sarah Menz, Environmental Administrator

Date Certified: 22-DEC-2023 14:02

A handwritten signature in black ink, appearing to be 'Sarah Menz', written in a cursive style.

Case Narrative

Unless otherwise noted, all samples included in this report were received in accordance with protocols referenced in Chapter 62-160, Florida Administrative Code (F.A.C.). Results published in this report pertain only to the samples as submitted to, and received by the laboratory. All times in this report are adjusted to the applicable Eastern Time Zone (EST or EDT).

Results for the following analytical groups are included in this report: Chlorophyll/Grain Size/BOD, Molecular and Overflow.

Scientific notation may be used in reporting very large or small values. Values reported using scientific notation will take the form of the following example: 1.3E+03, which is equivalent to 1.3×10^3 or 1300.

Unless otherwise noted, analytical values for soil and sediment samples are reported on a dry weight basis, and analytical values for waste and tissue samples are reported on a wet weight basis.

Results for TNI accredited tests met requirements established by The NELAC Institute. A double asterisk (**) is used to indicate an analyte/matrix/method for which the laboratory is not TNI accredited by the Florida Department of Health Environmental Laboratory Certification Program or where accreditation for that field of testing is not applicable.

Any significant anomalies or deviations from established protocols are documented in Non-Conformance Reports, which, where appropriate, are included within this analytical report. Additional comments related to specific analytical tests may be included as remarks following the analytical results for each sample. Such comments and remarks are for informational purposes only and are not intended to convey judgement about the usability of the reported data.

A quality control report on the performance of the test method for the submitted samples is included. Uncertainty associated with the analytical results contained in this report can be estimated from the reported quality assurance results and from published quality control acceptance limits for each analytical test. Matrix quality control results (matrix spike recoveries and matrix sample precision) pertain only to the matrix sample tested and do not necessarily reflect test method performance for other samples.

Typical matrix quality control (QC) measurements may include matrix spike recovery, matrix spike duplicate recovery, matrix spike precision and matrix sample precision. Not all matrix QC results may be available or reportable; where they are not an explanation is provided. Typical reasons for unavailable QC results include, but are not limited to, a) insufficient matrix sample to perform some or all QC measurements; b) analyte concentration in the sample replicated was too low for a meaningful measurement of precision and c) analyte concentration in the matrix sample spiked was too high (relative to the amount of analyte spiked) for a meaningful measurement of recovery. Where matrix QC results are unavailable, other method performance metrics (e.g., LCS recovery, LCS precision, surrogate recovery) may be used to assess performance of the method. Comments explaining any missing QC measurements are not intended to convey any adverse conclusions about the quality of the reported data.

Precision is reported as relative percent difference unless otherwise noted.

Quality Control codes as defined below may be used in this report to indicate results that are associated with one or more quality control elements which did not fall within established test method criteria. Such results may be qualified as estimates using a J qualifier as required by 62-160 F.A.C. Explanations are included in the report for any results that were reported as estimates for other reasons.

QC Codes used in this report may include:

- LCS – Recovery for the batch Laboratory Control Sample (LCS) was outside existing control limits;
- MS – Recovery for the batch matrix spike (MS) was outside existing control limits;
- CCV – Recovery for a continuing calibration verification (CCV) standard was outside existing control limits;
- SUR – Recovery of a surrogate (SUR) for associated analytes was outside existing control limits;
- RPD – The precision, measured as relative percent difference (RPD), of batch replicate measurements was outside existing control limits;
- RSD – The precision, measured as relative standard deviation (RSD), of batch replicate measurements was outside existing control limits;
- SMP – Sample - used precision derived from replicate analyses of a sample;

The following data qualifiers are used, where applicable, in this report as specified in 62-160 F.A.C.

- A - Value reported is the mean of two or more determinations.
- B - Results based on colony counts outside the acceptable range.
- I - The reported value is between the laboratory method detection limit and the laboratory practical quantitation limit.
- J - Estimated value and/or the analysis did not meet established quality control criteria.
- K - Actual value is known to be less than value given.
- L - Actual value is known to be greater than value given.
- N - Presumptive evidence of presence of material.
- O - Sampled, but analysis lost or not performed.
- Q - Sample held beyond normal holding time.
- T - Value reported is less than the criterion of detection.
- U - Material was analyzed for but not detected. The reported value is the method detection limit for the sample analyzed.
- V - Analyte was detected in both sample and method blank.
- X - Too few individuals to calculate SCI value.
- Y - The laboratory analysis was from an unpreserved or improperly preserved sample. The data may not be accurate.
- Z - Colonies were too numerous to count (TNTC).

Quality control information from overflow laboratories may not be included in this report. Please refer to the associated report from the overflow laboratory for additional information.

Sample Location: UNNAMED BRANCH 05 MI ABOVE TOMOKA RIVER

Collection Date/Time: 11/15/2023 11:20

Field ID: G5NE0609

Matrix: W-SURF-SLT

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2451954	SM 10150 B	Chlorophyll-a, Corrected	1.4	I	ug/L	P438610	
		Phaeophytin-a	0.90	U	ug/L	P438610	
		Chlorophyll-a, Uncorrected	1.6	I	ug/L	P438610	
2451957	Enterolert/QT	Enterococci-Quanti-Tray	86.5		MPN/100 mL		

Ref. Method and Comment:

SM 10150 B: Precision data is not available for at least one component due to the small amount of analyte in the QC sample. Refer to QA report for available precision data.

Sample Location: GROOVER BRANCH AT AIRPORT ROAD BRIDGE

Collection Date/Time: 11/15/2023 11:45

Field ID: G5NE0610

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2451955	SM 10150 B	Chlorophyll-a, Corrected	1.6	U	ug/L	P438610	
		Phaeophytin-a	1.8	U	ug/L	P438610	
		Chlorophyll-a, Uncorrected	1.2	U	ug/L	P438610	
2451958	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	622.1		MPN/100 mL		
2451959	SOP-PCR-1.0	HF183-qPCR**	3.3E+03	U	TSC/100 mL	P437732	
	SOP-PCR-1.4	BacR-qPCR**	1.4E+04	I	TSC/100 mL	P437732	
	SOP-PCR-1.5	DG3-qPCR**	2.6E+03	U	TSC/100 mL	P437732	

Ref. Method and Comment:

SM 10150 B: Lab dup result is 1.6 U for Chl-a Corrected, 1.2 U for Chl-a Uncorrected, and 1.8 U for Phaeophytin-a. Precision data is unavailable due to the small amount of analyte in the QC sample.

Sample Location: LAKE EATON CENTER

Collection Date/Time: 11/15/2023 13:10

Field ID: G1NE0954

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2451956	SM 10150 B	Chlorophyll-a, Corrected	1.1	I	ug/L	P438614	
		Phaeophytin-a	1.1	I	ug/L	P438614	
		Chlorophyll-a, Uncorrected	1.8		ug/L	P438614	

Ref. Method and Comment:

SM 10150 B: Precision data is not available for at least one component due to the small amount of analyte in the QC sample. Refer to QA report for available precision data.

Quality Assurance Report Method Blank Results

Reference Method: SM 10150 B

Batch ID: P438610

Component	Result	Code	Units
Chlorophyll-a, Corrected	0.82	U	ug/L
Chlorophyll-a, Uncorrected	0.60	U	ug/L
Phaeophytin-a	0.90	U	ug/L

Quality Assurance Report Method Blank Results

Reference Method: SM 10150 B
Batch ID: P438614

Component	Result	Code	Units
Chlorophyll-a, Corrected	0.82	U	ug/L
Chlorophyll-a, Uncorrected	0.60	U	ug/L
Phaeophytin-a	0.90	U	ug/L

Reference Method: SOP-PCR-1.0
Batch ID: P437732

Component	Result	Code	Units
HF183-qPCR	3.3E+03	U	TSC/100 mL

Reference Method: SOP-PCR-1.4
Batch ID: P437732

Component	Result	Code	Units
BacR-qPCR	3.0E+03	U	TSC/100 mL

Reference Method: SOP-PCR-1.5
Batch ID: P437732

Component	Result	Code	Units
DG3-qPCR	2.6E+03	U	TSC/100 mL

Quality Assurance Report Laboratory Control Sample Accuracy

Reference Method: SM 10150 B
Batch ID: P438610

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	89.4		P	84 - 116

Reference Method: SM 10150 B
Batch ID: P438614

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	91.0		P	84 - 116

Reference Method: SOP-PCR-1.0
Batch ID: P437732

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	94.1	71.1	P/P	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P437732

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
BacR-qPCR	98.8	68.5	P/P	50 - 175

Quality Assurance Report
Laboratory Control Sample Accuracy

Reference Method: SOP-PCR-1.5
Batch ID: P437732

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
DG3-qPCR	105	70.5	P/P	50 - 175

Quality Assurance Report
Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
Batch ID: P437732

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2451153	HF183-qPCR	88.4	92.4	P/P	50 - 175
2451192	HF183-qPCR	89.1		P	50 - 175
2451205	HF183-qPCR	100		P	50 - 175
2451206	HF183-qPCR	108		P	50 - 175
2451418	HF183-qPCR	81.5		P	50 - 175
2451419	HF183-qPCR	89.0		P	50 - 175
2451681	HF183-qPCR	85.4		P	50 - 175
2451959	HF183-qPCR	88.7		P	50 - 175
2451985	HF183-qPCR	84.9		P	50 - 175
2452066	HF183-qPCR	75.7		P	50 - 175
2452125	HF183-qPCR	105		P	50 - 175
2452126	HF183-qPCR	88.6		P	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P437732

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2451153	BacR-qPCR	84.0	89.9	P/P	50 - 175
2451192	BacR-qPCR	95.8		P	50 - 175
2451205	BacR-qPCR	91.8		P	50 - 175
2451206	BacR-qPCR	83.9		P	50 - 175
2451418	BacR-qPCR	64.1		P	50 - 175
2451419	BacR-qPCR	84.6		P	50 - 175
2451681	BacR-qPCR	85.3		P	50 - 175
2451959	BacR-qPCR	78.4		P	50 - 175
2451985	BacR-qPCR	81.0		P	50 - 175
2452066	BacR-qPCR	84.4		P	50 - 175
2452125	BacR-qPCR	81.3		P	50 - 175
2452126	BacR-qPCR	62.8		P	50 - 175

Reference Method: SOP-PCR-1.5
Batch ID: P437732

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2451153	DG3-qPCR	94.4	100	P/P	50 - 175
2451192	DG3-qPCR	100		P	50 - 175
2451205	DG3-qPCR	106		P	50 - 175
2451206	DG3-qPCR	105		P	50 - 175
2451418	DG3-qPCR	82.5		P	50 - 175
2451419	DG3-qPCR	94.9		P	50 - 175
2451681	DG3-qPCR	99.4		P	50 - 175
2451959	DG3-qPCR	105		P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.5
Batch ID: P437732

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2451985	DG3-qPCR	104		P	50 - 175
2452066	DG3-qPCR	96.9		P	50 - 175
2452125	DG3-qPCR	91.8		P	50 - 175
2452126	DG3-qPCR	60.6		P	50 - 175

Quality Assurance Report Precision

Reference Method: SOP-PCR-1.0
Batch ID: P437732

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2451153	HF183-qPCR	4.40	Spike	P	0 - 25

Reference Method: SOP-PCR-1.4
Batch ID: P437732

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2451153	BacR-qPCR	6.81	Spike	P	0 - 25

Reference Method: SOP-PCR-1.5
Batch ID: P437732

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2451153	DG3-qPCR	5.88	Spike	P	0 - 25

* Sample, spike and/or laboratory control sample precision (LCS) is reported.

Quality Assurance Report Surrogates

Lab Sample ID: 2451959
Field Sample ID: G5NE0610

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	75.0	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	89.3	P	50 - 300
SOP-PCR-1.5	SKETA-qPCR	78.5	P	50 - 300

Quality Assurance Report Summary

Ref. Method	Analyte	LCS % Recovery	MS % Recovery	LCS	Precision SMP	MS
SM 10150 B	Chlorophyll-a, Corrected	89.4				
	Chlorophyll-a, Corrected	91.0				

Quality Assurance Report
Summary

Ref. Method	Analyte	LCS % Recovery		MS % Recovery			Precision SMP	MS
SOP-PCR-1.0	HF183-qPCR	71.1	94.1	88.4	92.4	89.1		4.40
SOP-PCR-1.4	BacR-qPCR	68.5	98.8	84.0	89.9	95.8		6.81
SOP-PCR-1.5	DG3-qPCR	70.5	105	94.4	100	100		5.88

Reference Method Descriptions

Method	Description	Associated Samples
Enterolert/QT	Enterococci by Enterolert/QT by Diversified Environmental Laboratories in Jacksonville	2451957
SM 10150 B	Phytoplankton chlorophyll-a corrected, uncorrected, and phaeophytin by spectrophotometry	2451954, 2451955, 2451956
SM 9223 B Quanti-Tray	E. coli by Colilert/QT by Diversified Environmental Laboratories in Jacksonville	2451958
SOP-PCR-1.0	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	2451959
SOP-PCR-1.4	Detection and quantification of Ruminant specific Bacteroidetes BacR genetic marker with quantitative polymerase chain reaction	2451959
SOP-PCR-1.5	Detection and quantification of canine-specific DG3 Bacteroides genetic marker with quantitative polymerase chain reaction	2451959

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
Enterolert/QT	11/15/2023	11/15/2023 15:33		11/16/2023 15:55	Diversified-Jacksonville	2451957
SM 10150 B	11/16/2023	11/16/2023 13:59	Tynan A Branch	11/29/2023 16:10	Tynan A Branch	2451954, 2451955
	11/16/2023	11/16/2023 13:59	Tynan A Branch	11/30/2023 09:08	Callie Williams	2451956
SM 9223 B Quanti-Tray	11/15/2023	11/15/2023 15:44		11/16/2023 10:04	Diversified-Jacksonville	2451958
SOP-PCR-1.0	11/16/2023	11/16/2023 13:15	Madeline Gruver	12/05/2023 08:47	Roberta Tatti	2451959
SOP-PCR-1.4	11/16/2023	11/16/2023 13:15	Madeline Gruver	12/06/2023 08:50	Roberta Tatti	2451959
SOP-PCR-1.5	11/16/2023	11/16/2023 13:15	Madeline Gruver	12/06/2023 09:50	Roberta Tatti	2451959