

Biological Analysis Report

NE-DIST-2018-09-11-02

Florida Department of Environmental Protection
Central Laboratory
2600 Blair Stone Road
Tallahassee, FL 32399-2400
DOH Accreditation E31780

Overflow Analyses Performed By:
Advanced Environmental Laboratories - Gainesville
6815 SW Archer Road
Gainesville, Florida 32608
DOH Accreditation E82001

Event Description: **Alachua County Sampling**
Request ID: **RQ-2018-09-10-56**
Customer: **NE-DIST**
Project ID: **SMP**

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

For additional information please contact
Cheryl Swanson - Administrator
J. Marshall Faircloth - Aquatic Toxicology
Jacqueline Savage - Bench Biology
Jacqueline Savage - Microbiology
Puja Jasrotia, Ph.D. - Molecular Biology
Amelia Rankin - Taxonomy (Invertebrate & Algal)
Thekkekalathil Chandrasekhar, Ph.D, QA Officer
Phone (850) 245-8177

Certified by: Cheryl A. Swanson, Program Administrator

Date Certified: 17-DEC-2018 17:39



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 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: MILL CREEK - E OF OLD BELAMY RD

Collection Date/Time: 09/10/2018 10:50

Field ID: G1NE0907

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2023634	SM 10200 H (mod.)	Chlorophyll-a, Corrected	0.55	U	ug/L	P351959	
		Phaeophytin-a	0.40	U	ug/L	P351959	
		Chlorophyll-a, Uncorrected	0.61	I	ug/L	P351959	
2023638	EPA 1603	Escherichia coli-Membrane Filter	810	B	cfu/100 mL		
2023642	SOP-PCR-1.0	HF183-qPCR**	200	I	GEU/100 mL	P353497	LCS
	SOP-PCR-4.0	HF183 - Human Source Marker**	Above Threshold				

Sample Location: TURKEY CREEK UPSTREAM OF THE BRIDGE ON NW 74TH TERR

Collection Date/Time: 09/10/2018 11:30

Field ID: G1NE0915

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2023635	SM 10200 H (mod.)	Chlorophyll-a, Corrected	1.4	I	ug/L	P351959	
		Phaeophytin-a	0.67	I	ug/L	P351959	
		Chlorophyll-a, Uncorrected	1.8		ug/L	P351959	
2023639	EPA 1603	Escherichia coli-Membrane Filter	560		cfu/100 mL		
2023643	SOP-PCR-1.0	HF183-qPCR**	500	I	GEU/100 mL	P353497	LCS
	SOP-PCR-4.0	HF183 - Human Source Marker**	Above Threshold				

Sample Location: POSSUM CREEK - S OF NW 16TH AVE

Collection Date/Time: 09/10/2018 12:45

Field ID: G1NE0900

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2023636	SM 10200 H (mod.)	Chlorophyll-a, Corrected	0.55	U	ug/L	P351959	
		Phaeophytin-a	0.40	U	ug/L	P351959	
		Chlorophyll-a, Uncorrected	0.40	U	ug/L	P351959	
2023640	EPA 1603	Escherichia coli-Membrane Filter	660		cfu/100 mL		
2023644	SOP-PCR-1.0	HF183-qPCR**	1.27E+03		GEU/100 mL	P353497	LCS
	SOP-PCR-4.0	HF183 - Human Source Marker**	Above Threshold				

Sample Location: HOGTOWN CREEK - RING PARK, N OF NW 16TH AVE (BETWEEN NW 18TH)

Collection Date/Time: 09/10/2018 13:20

Field ID: G1NE0912

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2023637	SM 10200 H (mod.)	Chlorophyll-a, Corrected	0.55	U	ug/L	P351959	
		Phaeophytin-a	0.40	U	ug/L	P351959	
		Chlorophyll-a, Uncorrected	0.52	I	ug/L	P351959	
2023641	EPA 1603	Escherichia coli-Membrane Filter	880	B	cfu/100 mL		
2023645	SOP-PCR-1.0	HF183-qPCR**	300	I	GEU/100 mL	P353497	LCS
	SOP-PCR-4.0	HF183 - Human Source Marker**	Above Threshold				

Quality Assurance Report

Method Blank Results

Quality Assurance Report Method Blank Results

Reference Method: SM 10200 H (mod.)
Batch ID: P351959

Component	Result	Code	Units
Chlorophyll-a, Corrected	0.55	U	ug/L
Chlorophyll-a, Uncorrected	0.40	U	ug/L
Phaeophytin-a	0.40	U	ug/L

Reference Method: SOP-PCR-1.0
Batch ID: P353497

Component	Result	Code	Units
HF183-qPCR	280	U	GEU/100 mL

Quality Assurance Report Laboratory Control Sample Accuracy

Reference Method: SM 10200 H (mod.)
Batch ID: P351959

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	87.7		P	84 - 116

Reference Method: SOP-PCR-1.0
Batch ID: P353497

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	108	187	P/F	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
Batch ID: P353497

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2023642	HF183-qPCR	85.1		P	50 - 175
2023643	HF183-qPCR	85.9		P	50 - 175
2023644	HF183-qPCR	106		P	50 - 175
2023645	HF183-qPCR	95.6		P	50 - 175
2026896	HF183-qPCR	68.3		P	50 - 175
2026986	HF183-qPCR	92.9		P	50 - 175
2026987	HF183-qPCR	61.3	54.7	P/P	50 - 175
2026988	HF183-qPCR	78.4		P	50 - 175
2027411	HF183-qPCR	90.0		P	50 - 175
2027445	HF183-qPCR	79.0		P	50 - 175
2030524	HF183-qPCR	94.6		P	50 - 175
2034865	HF183-qPCR	83.6		P	50 - 175

Quality Assurance Report Precision

Quality Assurance Report Precision

Reference Method: SM 10200 H (mod.)
 Batch ID: P351959

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2023538	Chlorophyll-a, Corrected	4.24	Sample	P	0 - 25
2023538	Chlorophyll-a, Uncorrected	4.48	Sample	P	0 - 25
2023538	Phaeophytin-a	6.91	Sample	P	0 - 25

Reference Method: SOP-PCR-1.0
 Batch ID: P353497

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2026987	HF183-qPCR	11.3	Spike	P	0 - 25

* Sample, spike and/or laboratory control sample precision (LCS) is reported.
 Replicate spike precision may be reported when sample results are below quantifiable levels.

Quality Assurance Report Surrogates

Lab Sample ID: 2023642
 Field Sample ID: G1NE0907

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	136	P	50 - 300

Lab Sample ID: 2023643
 Field Sample ID: G1NE0915

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	127	P	50 - 300

Lab Sample ID: 2023644
 Field Sample ID: G1NE0900

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	143	P	50 - 300

Lab Sample ID: 2023645
 Field Sample ID: G1NE0912

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	141	P	50 - 300

Quality Assurance Report Summary

Ref. Method	Analyte	LCS % Recovery	MS % Recovery	LCS	Precision SMP	MS
SM 10200 H (mod.)	Chlorophyll-a, Corrected	87.7			4.24	
	Chlorophyll-a, Uncorrected				4.48	
	Phaeophytin-a				6.91	

Quality Assurance Report Summary

Ref. Method	Analyte	LCS % Recovery		MS % Recovery			LCS	Precision SMP	MS
SOP-PCR-1.0	HF183-qPCR	187	108	85.1	85.9	106			11.3
				95.6					

Reference Method Descriptions

Method / DoH Cert #	Description	<u>Associated Samples</u>
EPA 1603 / E82001	Escherichia coli by membrane filter method by Advanced Environmental Laboratory in Gainesville	2023638, 2023639, 2023640, 2023641
SM 10200 H (mod.) / E31780	Phytoplankton chlorophyll-a corrected, uncorrected, and phaeophytin by spectrophotometry	2023634, 2023635, 2023636, 2023637
SOP-PCR-1.0 / E31780	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	2023642, 2023643, 2023644, 2023645
SOP-PCR-4.0 / E31780	Preparation of Samples by Filtration and held for qPCR Analysis	2023642, 2023643, 2023644, 2023645

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
EPA 1603	09/10/2018	09/10/2018 16:07		09/11/2018 07:14	AEL - Gainesville	2023638, 2023639, 2023640, 2023641
SM 10200 H (mod.)	09/11/2018	09/11/2018 14:43	Mariam Hanna	09/19/2018 10:40	Peterson Janvier	2023634, 2023635, 2023636, 2023637
SOP-PCR-1.0	09/11/2018	09/11/2018 14:15	Sarah Menz	10/24/2018 14:19	Mailin Lopez	2023642, 2023643, 2023644, 2023645
SOP-PCR-4.0	09/11/2018	09/11/2018 14:15	Angela Smith	10/23/2018 09:59	Sarah Menz	2023642, 2023643, 2023644, 2023645