

Biological Analysis Report

NE-DIST-2017-11-01-01

Florida Department of Environmental Protection
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DOH Accreditation E31780

Overflow Analyses Performed By:
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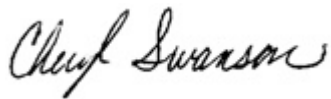
Event Description: **Gainesville 2**
Request ID: **RQ-2017-10-30-10**
Customer: **NE-DIST**
Project ID: **SMP**

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Date Certified: 24-JUL-2018 18:22



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: Hatchett Crk @ SR 26

Collection Date/Time: 10/31/2017 09:40

Field ID: G1NE0096 10312017

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1941745	SM 10200 H (mod.)	Chlorophyll-a, Corrected	0.55	U	ug/L	P335006	
		Phaeophytin-a	0.40	U	ug/L	P335006	
		Chlorophyll-a, Uncorrected	0.40	U	ug/L	P335006	
1941746	EPA 1603	Escherichia coli-Membrane Filter	20	U	cfu/100 mL		
1941755	SOP-PCR-1.0	HF183-qPCR**	280	U	GEU/100 mL	P334516	
	SOP-PCR-4.0	HF183 - Human Source Marker**	Below Threshold				

Ref. Method and Comment:

SM 10200 H (mod.): 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.

SOP-PCR-1.0: No amplification of HF183 marker.

Sample Location: Lake Forest Crk @ University Ave

Collection Date/Time: 10/31/2017 10:05

Field ID: G1NE0098 10272017

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1941742	SM 10200 H (mod.)	Chlorophyll-a, Corrected	1.8		ug/L	P334832	
		Phaeophytin-a	0.40	U	ug/L	P334832	
		Chlorophyll-a, Uncorrected	1.7	A	ug/L	P334832	

Ref. Method and Comment:

SM 10200 H (mod.): Laboratory duplicate result is 1.6 I for Chlorophyll-a Corrected and 0.40 U for Phaeophytin-a. Precision data is unavailable due to the small amount of analyte in the QC sample.

Sample Location: Lake Forest Crk @ University Ave

Collection Date/Time: 10/31/2017 10:05

Field ID: G1NE0098 10272017

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1941747	EPA 1603	Escherichia coli-Membrane Filter	710		cfu/100 mL		
1941759	SOP-PCR-1.0	HF183-qPCR**	26	T	GEU/100 mL	P334516	
	SOP-PCR-4.0	HF183 - Human Source Marker**	Above Threshold				

Sample Location: Sweetwater Creek above wetland

Collection Date/Time: 10/31/2017 11:15

Field ID: G1NE0001 10312017

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1941748	EPA 1603	Escherichia coli-Membrane Filter	10	B	cfu/100 mL		
1941760	SOP-PCR-1.0	HF183-qPCR**	35	T	GEU/100 mL	P334516	
	SOP-PCR-4.0	HF183 - Human Source Marker**	Below Threshold				

Sample Location: Bivens Arm Outlet @ SR 331

Collection Date/Time: 10/31/2017 11:25

Field ID: G1NE0208 10312017

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1941743	SM 10200 H (mod.)	Chlorophyll-a, Corrected	1.5	I	ug/L	P334832	
		Phaeophytin-a	0.40	U	ug/L	P334832	
		Chlorophyll-a, Uncorrected	1.6		ug/L	P334832	

Ref. Method and Comment:

SM 10200 H (mod.): 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: Lake Kanapaha 300m N of 56th

Collection Date/Time: 10/31/2017 11:50

Field ID: G2NE0933 10312017

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1941744	SM 10200 H (mod.)	Chlorophyll-a, Corrected	3.6		ug/L	P334832	
		Phaeophytin-a	1.1	I	ug/L	P334832	
		Chlorophyll-a, Uncorrected	4.2		ug/L	P334832	

Ref. Method and Comment:

SM 10200 H (mod.): 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 10200 H (mod.)

Batch ID: P334832

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: SM 10200 H (mod.)

Batch ID: P335006

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: P334516

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

Quality Assurance Report Laboratory Control Sample Accuracy

Reference Method: SM 10200 H (mod.)

Batch ID: P334832

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	103		P	84 - 116

Reference Method: SM 10200 H (mod.)

Batch ID: P335006

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	107		P	84 - 116

Reference Method: SOP-PCR-1.0

Batch ID: P334516

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	109	112	P/P	50 - 175

Quality Assurance Report Laboratory Control Sample Accuracy

Reference Method: SOP-PCR-1.0
 Batch ID: P334516

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	120	95.7	P/P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
 Batch ID: P334516

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
1941755	HF183-qPCR	75.6	82.4	P/P	50 - 175
1941759	HF183-qPCR	71.7		P	50 - 175
1941760	HF183-qPCR	88.8		P	50 - 175
1941930	HF183-qPCR	65.3		P	50 - 175
1941931	HF183-qPCR	74.5		P	50 - 175
1942101	HF183-qPCR	79.5		P	50 - 175
1942102	HF183-qPCR	86.8		P	50 - 175
1942368	HF183-qPCR	85.0		P	50 - 175
1942369	HF183-qPCR	86.2		P	50 - 175
1942370	HF183-qPCR	63.1		P	50 - 175
1942371	HF183-qPCR	60.3		P	50 - 175
1942489	HF183-qPCR	73.2		P	50 - 175
1942490	HF183-qPCR	75.8		P	50 - 175

Quality Assurance Report Precision

Reference Method: SM 10200 H (mod.)
 Batch ID: P334832

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
1941742	Chlorophyll-a, Uncorrected	11.2	Sample	P	0 - 25

Reference Method: SOP-PCR-1.0
 Batch ID: P334516

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
1941755	HF183-qPCR	8.71	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: 1941755
 Field Sample ID: G1NE0096 10312017

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
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Quality Assurance Report Surrogates

Lab Sample ID: 1941755
 Field Sample ID: G1NE0096 10312017

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	92.0	P	50 - 300

Lab Sample ID: 1941759
 Field Sample ID: G1NE0098 10272017

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	90.4	P	50 - 300

Lab Sample ID: 1941760
 Field Sample ID: G1NE0001 10312017

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	91.2	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	103								
	Chlorophyll-a, Corrected	107								
	Chlorophyll-a, Uncorrected								11.2	
SOP-PCR-1.0	HF183-qPCR	109	95.7	120	79.5	75.6	82.4			8.71
		112			71.7					

Reference Method Descriptions

Method / DoH Cert #	Description	Associated Samples
EPA 1603 / E82001	Escherichia coli by membrane filter method using modified mTEC by Advanced Environmental Gainesville Overflow Laboratories for NED	1941746, 1941747, 1941748
SM 10200 H (mod.) / E31780	Phytoplankton chlorophyll-a corrected, uncorrected, and phaeophytin by spectrophotometry	1941742, 1941743, 1941744, 1941745
SOP-PCR-1.0 / E31780	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	1941755, 1941759, 1941760
SOP-PCR-4.0 / E31780	Preparation of Samples by Filtration and held for qPCR Analysis	1941755, 1941759, 1941760

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
EPA 1603	10/31/2017	10/31/2017 17:35		11/01/2017 19:10	AEL - Gainesville	1941746, 1941747, 1941748
SM 10200 H (mod.)	11/01/2017	11/01/2017 13:04	Cathy Sherrer	11/07/2017 08:05	Kim M. Spencer	1941742, 1941743, 1941744
	11/01/2017	11/01/2017 13:04	Cathy Sherrer	11/08/2017 13:50	Kim M. Spencer	1941745
SOP-PCR-1.0	11/01/2017	11/01/2017 13:30	Avril Wood	11/08/2017 11:48	Sarah Menz	1941755, 1941759, 1941760
SOP-PCR-4.0	11/01/2017	11/01/2017 13:30	Avril Wood	12/21/2017 13:17	Puja Jasrotia	1941755, 1941759, 1941760