

Biological Analysis Report

NE-DIST-2018-06-15-03

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-2018-06-11-27**
Customer: **NE-DIST**
Project ID: **SMP**

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Date Certified: 20-JUL-2018 09:45



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: 06/14/2018 11:00

Field ID: G1NE0907

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2001293	SM 10200 H (mod.)	Chlorophyll-a, Corrected	0.91	I	ug/L	P347381	
		Phaeophytin-a	0.48	I	ug/L	P347381	
		Chlorophyll-a, Uncorrected	1.2		ug/L	P347381	
2001296	EPA 1603	Escherichia coli-Membrane Filter	730		cfu/100 mL		
2001299	SOP-PCR-1.0	HF183-qPCR**	200	I	GEU/100 mL	P347849	
	SOP-PCR-4.0	HF183 - Human Source Marker**	Above Threshold				

Sample Location: Paraners Branch

Collection Date/Time: 06/14/2018 13:45

Field ID: G1NE0917

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2001294	SM 10200 H (mod.)	Chlorophyll-a, Corrected	2.8		ug/L	P347381	
		Phaeophytin-a	16		ug/L	P347381	
		Chlorophyll-a, Uncorrected	12		ug/L	P347381	
2001297	EPA 1603	Escherichia coli-Membrane Filter	870	B	cfu/100 mL		
2001300	SOP-PCR-1.0	HF183-qPCR**	140	U	GEU/100 mL	P348050	
	SOP-PCR-1.4	BacR-qPCR**	3.28E+08	J	TSC/100 mL	P348050	LCS, MS
	SOP-PCR-4.0	HF183 - Human Source Marker**	Above Threshold				

Ref. Method and Comment:

SOP-PCR-1.4: Quality control failure(s) observed. See QA report for details.

Sample Location: Paraners Branch - E of CR 1491

Collection Date/Time: 06/14/2018 12:30

Field ID: G1NE0906

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2001295	SM 10200 H (mod.)	Chlorophyll-a, Corrected	2.2		ug/L	P347381	
		Phaeophytin-a	3.8		ug/L	P347381	
		Chlorophyll-a, Uncorrected	4.6		ug/L	P347381	
2001298	EPA 1603	Escherichia coli-Membrane Filter	2800		cfu/100 mL		
2001301	SOP-PCR-1.0	HF183-qPCR**	140	U	GEU/100 mL	P348050	
	SOP-PCR-1.4	BacR-qPCR**	6.23E+06		TSC/100 mL	P348050	LCS, MS
	SOP-PCR-4.0	HF183 - Human Source Marker**	Above Threshold				

Quality Assurance Report Method Blank Results

Reference Method: SM 10200 H (mod.)

Batch ID: P347381

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: P347849

Component	Result	Code	Units
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Quality Assurance Report Method Blank Results

Reference Method: SOP-PCR-1.0
Batch ID: P347849

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

Reference Method: SOP-PCR-1.0
Batch ID: P348050

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

Reference Method: SOP-PCR-1.4
Batch ID: P348050

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

Quality Assurance Report Laboratory Control Sample Accuracy

Reference Method: SM 10200 H (mod.)
Batch ID: P347381

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	94.1		P	84 - 116

Reference Method: SOP-PCR-1.0
Batch ID: P347849

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	93.3	133	P/P	50 - 175

Reference Method: SOP-PCR-1.0
Batch ID: P348050

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	106	158	P/P	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P348050

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
BacR-qPCR	120	213	P/F	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
Batch ID: P347849

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
1996396	HF183-qPCR	80.1		P	50 - 175
1999617	HF183-qPCR	93.1		P	50 - 175
2000024	HF183-qPCR	81.5		P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
Batch ID: P347849

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2000025	HF183-qPCR	87.0		P	50 - 175
2000026	HF183-qPCR	87.1		P	50 - 175
2000997	HF183-qPCR	77.4		P	50 - 175
2001000	HF183-qPCR	102		P	50 - 175
2001245	HF183-qPCR	92.0		P	50 - 175
2001299	HF183-qPCR	83.1		P	50 - 175
2004524	HF183-qPCR	94.3	92.3	P/P	50 - 175
2005183	HF183-qPCR	62.1		P	50 - 175
2005186	HF183-qPCR	75.2		P	50 - 175

Reference Method: SOP-PCR-1.0
Batch ID: P348050

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2001300	HF183-qPCR	84.6		P	50 - 175
2001301	HF183-qPCR	98.9		P	50 - 175
2001768	HF183-qPCR	99.2		P	50 - 175
2002640	HF183-qPCR	64.2		P	50 - 175
2002641	HF183-qPCR	94.2		P	50 - 175
2002645	HF183-qPCR	84.8		P	50 - 175
2005554	HF183-qPCR	106		P	50 - 175
2005555	HF183-qPCR	81.7		P	50 - 175
2005556	HF183-qPCR	99.9		P	50 - 175
2005557	HF183-qPCR	106		P	50 - 175
2005771	HF183-qPCR	100		P	50 - 175
2005772	HF183-qPCR	95.0		P	50 - 175
2005773	HF183-qPCR	68.1	68.9	P/P	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P348050

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2001300	BacR-qPCR	-91.4		F	50 - 175
2001301	BacR-qPCR	76.3		P	50 - 175
2002640	BacR-qPCR	4.98		F	50 - 175
2002641	BacR-qPCR	78.3		P	50 - 175
2002645	BacR-qPCR	86.8		P	50 - 175
2005554	BacR-qPCR	96.9		P	50 - 175
2005555	BacR-qPCR	64.1		P	50 - 175
2005556	BacR-qPCR	84.8		P	50 - 175
2005557	BacR-qPCR	129		P	50 - 175
2005771	BacR-qPCR	96.5		P	50 - 175
2005772	BacR-qPCR	78.2		P	50 - 175
2005773	BacR-qPCR	69.3	69.5	P/P	50 - 175

Quality Assurance Report Precision

Quality Assurance Report Precision

Reference Method: SM 10200 H (mod.)
Batch ID: P347381

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2001311	Chlorophyll-a, Corrected	9.04	Sample	P	0 - 25
2001311	Chlorophyll-a, Uncorrected	8.70	Sample	P	0 - 25
2001311	Phaeophytin-a	2.41	Sample	P	0 - 25

Reference Method: SOP-PCR-1.0
Batch ID: P347849

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2004524	HF183-qPCR	2.16	Spike	P	0 - 25

Reference Method: SOP-PCR-1.0
Batch ID: P348050

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2005773	HF183-qPCR	1.27	Spike	P	0 - 25

Reference Method: SOP-PCR-1.4
Batch ID: P348050

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2005773	BacR-qPCR	0.195	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: 2001299
Field Sample ID: G1NE0907

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	115	P	50 - 300

Lab Sample ID: 2001300
Field Sample ID: G1NE0917

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	110	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	104	P	50 - 300

Lab Sample ID: 2001301
Field Sample ID: G1NE0906

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	109	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	101	P	50 - 300

Quality Assurance Report Summary

Ref. Method	Analyte	LCS % Recovery		MS % Recovery			LCS	Precision	MS
								SMP	
SM 10200 H (mod.)	Chlorophyll-a, Corrected	94.1						9.04	
	Chlorophyll-a, Uncorrected							8.70	
	Phaeophytin-a							2.41	
SOP-PCR-1.0	HF183-qPCR	133	93.3	94.3	92.3	92.0			2.16
				80.1					
	HF183-qPCR	158	106	98.9	99.2	64.2			1.27
SOP-PCR-1.4				94.2					
	BacR-qPCR	213	120	76.3	4.98	78.3			0.195
				86.8					

Reference Method Descriptions

Method / DoH Cert #	Description	Associated Samples
EPA 1603 / E82001	Escherichia coli by membrane filter method using modified mTEC by Advanced Environmental Laboratory-Gainesville	2001296, 2001297, 2001298
SM 10200 H (mod.) / E31780	Phytoplankton chlorophyll-a corrected, uncorrected, and phaeophytin by spectrophotometry	2001293, 2001294, 2001295
SOP-PCR-1.0 / E31780	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	2001299, 2001300, 2001301
SOP-PCR-1.4 / E31780	Detection and quantification of Ruminant specific Bacteroidetes BacR genetic marker with quantitative polymerase chain reaction	2001300, 2001301
SOP-PCR-4.0 / E31780	Preparation of Samples by Filtration and held for qPCR Analysis	2001299, 2001300, 2001301

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
EPA 1603	06/14/2018	06/14/2018 19:00		06/15/2018 18:10	AEL - Gainesville	2001296
	06/14/2018	06/14/2018 20:15		06/15/2018 18:25	AEL - Gainesville	2001297, 2001298
SM 10200 H (mod.)	06/15/2018	06/15/2018 13:25	Mariam Hanna	06/22/2018 09:00	Carina A. Tull	2001293, 2001294, 2001295
SOP-PCR-1.0	06/15/2018	06/15/2018 13:00	Angela Smith	07/09/2018 15:26	Mailin Lopez	2001299
	06/15/2018	06/15/2018 13:00	Puja Jasrotia	07/13/2018 10:26	Mailin Lopez	2001300, 2001301
SOP-PCR-1.4	06/15/2018	06/15/2018 13:00	Puja Jasrotia	07/16/2018 12:13	Sarah Menz	2001300, 2001301
SOP-PCR-4.0	06/15/2018	06/15/2018 13:00	Angela Smith	07/09/2018 09:36	Puja Jasrotia	2001299, 2001300, 2001301