

Biological Analysis Report

SW-DIST-2019-09-10-01

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

Overflow Analyses Performed By:
Advanced Environmental Laboratories - Tampa
9610 Princess Palm Avenue
Tampa, Florida 33619
DOH Accreditation E84589

Event Description: **SMP: Baker, Bellow Outlet, New River**
Request ID: **RQ-2019-09-09-08**
Customer: **SW-DIST**
Project ID: **SMP**

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Date Certified: 07-OCT-2019 12:54



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: Baker Creek @ Thonotosassa Rd

Collection Date/Time: 09/09/2019 10:20

Field ID: G2SW0012

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2117237	SM 10200 H (mod.)	Chlorophyll-a, Corrected	13		ug/L	P370557	
		Phaeophytin-a	3.5		ug/L	P370557	
		Chlorophyll-a, Uncorrected	15		ug/L	P370557	
2117244	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	160		MPN/100 mL		

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: Baker Creek @ Thonotosassa Rd

Collection Date/Time: 09/09/2019 10:25

Field ID: G2SW0012_DUP

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2117238	SM 10200 H (mod.)	Chlorophyll-a, Corrected	12		ug/L	P370557	
		Phaeophytin-a	3.6		ug/L	P370557	
		Chlorophyll-a, Uncorrected	15		ug/L	P370557	
2117245	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	176		MPN/100 mL		

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: Field_Blank

Collection Date/Time: 09/09/2019 10:30

Field ID: FB 09092019

Matrix: W-FIELD-BK

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2117239	SM 10200 H (mod.)	Chlorophyll-a, Corrected	0.62	U	ug/L	P370557	
		Phaeophytin-a	0.68	U	ug/L	P370557	
		Chlorophyll-a, Uncorrected	0.45	U	ug/L	P370557	
2117246	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	1	U	MPN/100 mL		

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: Bellows Lake Outlet

Collection Date/Time: 09/09/2019 11:10

Field ID: G1SW0060

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2117241	SM 10200 H (mod.)	Chlorophyll-a, Corrected	40		ug/L	P370559	
		Phaeophytin-a	14		ug/L	P370559	
		Chlorophyll-a, Uncorrected	50		ug/L	P370559	
2117243	SOP-PCR-1.0	HF183-qPCR**	6.74E+03		GEU/100 mL	P370137	LCS
2117248	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	2240		MPN/100 mL		

Sample Location: New River @ Morris Bridge Rd

Collection Date/Time: 09/09/2019 09:00

Field ID: G2SW0061

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2117240	SM 10200 H (mod.)	Chlorophyll-a, Corrected	2.1	U	ug/L	P370557	
		Phaeophytin-a	2.2	U	ug/L	P370557	

Field ID: G2SW0061

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2117240	SM 10200 H (mod.)	Chlorophyll-a, Uncorrected	2.7	I	ug/L	P370557	
2117242	SOP-PCR-1.0	HF183-qPCR**	140	U	GEU/100 mL	P370137	LCS
	SOP-PCR-1.4	BacR-qPCR**	7.40E+04		TSC/100 mL	P370137	
2117247	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	394		MPN/100 mL		

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: P370557

Component	Result	Code	Units
Chlorophyll-a, Corrected	0.62	U	ug/L
Chlorophyll-a, Uncorrected	0.45	U	ug/L
Phaeophytin-a	0.68	U	ug/L

Reference Method: SM 10200 H (mod.)

Batch ID: P370559

Component	Result	Code	Units
Chlorophyll-a, Corrected	0.62	U	ug/L
Chlorophyll-a, Uncorrected	0.45	U	ug/L
Phaeophytin-a	0.68	U	ug/L

Reference Method: SOP-PCR-1.0

Batch ID: P370137

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

Reference Method: SOP-PCR-1.4

Batch ID: P370137

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: P370557

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	88.8		P	84 - 116

Reference Method: SM 10200 H (mod.)

Batch ID: P370559

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	90.2		P	84 - 116

Quality Assurance Report Laboratory Control Sample Accuracy

Reference Method: SOP-PCR-1.0
Batch ID: P370137

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	126	274	P/F	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P370137

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
BacR-qPCR	106	143	P/P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
Batch ID: P370137

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2116409	HF183-qPCR	113		P	50 - 175
2116489	HF183-qPCR	120		P	50 - 175
2116490	HF183-qPCR	131	118	P/P	50 - 175
2116588	HF183-qPCR	95.9		P	50 - 175
2116702	HF183-qPCR	111		P	50 - 175
2116703	HF183-qPCR	102		P	50 - 175
2116704	HF183-qPCR	104		P	50 - 175
2116918	HF183-qPCR	93.6		P	50 - 175
2117242	HF183-qPCR	82.6		P	50 - 175
2117243	HF183-qPCR	68.4		P	50 - 175
2117278	HF183-qPCR	93.2		P	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P370137

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2116409	BacR-qPCR	97.8		P	50 - 175
2116489	BacR-qPCR	90.5		P	50 - 175
2116490	BacR-qPCR	90.8	87.0	P/P	50 - 175
2116588	BacR-qPCR	79.3		P	50 - 175
2116702	BacR-qPCR	99.1		P	50 - 175
2116703	BacR-qPCR	96.7		P	50 - 175
2116704	BacR-qPCR	80.3		P	50 - 175
2116918	BacR-qPCR	51.3		P	50 - 175
2117242	BacR-qPCR	51.7		P	50 - 175
2117278	BacR-qPCR	63.1		P	50 - 175

Quality Assurance Report Precision

Reference Method: SM 10200 H (mod.)
Batch ID: P370557

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2117113	Chlorophyll-a, Corrected	5.45	Sample	P	0 - 25

Quality Assurance Report Precision

Reference Method: SM 10200 H (mod.)
Batch ID: P370557

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2117113	Chlorophyll-a, Uncorrected	6.16	Sample	P	0 - 25

Reference Method: SM 10200 H (mod.)
Batch ID: P370559

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2117312	Chlorophyll-a, Corrected	5.83	Sample	P	0 - 25
2117312	Chlorophyll-a, Uncorrected	6.08	Sample	P	0 - 25
2117312	Phaeophytin-a	7.65	Sample	P	0 - 25

Reference Method: SOP-PCR-1.0
Batch ID: P370137

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2116490	HF183-qPCR	10.5	Spike	P	0 - 25

Reference Method: SOP-PCR-1.4
Batch ID: P370137

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2116490	BacR-qPCR	4.35	Spike	P	0 - 25

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

Quality Assurance Report Surrogates

Lab Sample ID: 2117242
Field Sample ID: G2SW0061

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	74.6	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	81.0	P	50 - 300

Lab Sample ID: 2117243
Field Sample ID: G1SW0060

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	68.8	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	88.8			5.45	

Quality Assurance Report Summary

Ref. Method	Analyte	LCS % Recovery		MS % Recovery			LCS	Precision	MS
								SMP	
SM 10200 H (mod.)	Chlorophyll-a, Corrected	90.2						5.83	
	Chlorophyll-a, Uncorrected							6.16	
	Chlorophyll-a, Uncorrected							6.08	
	Phaeophytin-a							7.65	
SOP-PCR-1.0	HF183-qPCR	274	126	113	120	131			10.5
SOP-PCR-1.4	BacR-qPCR	143 106		118					4.35
				97.8	90.5	90.8			
				87.0					

Reference Method Descriptions

Method / DoH Cert #	Description	Associated Samples
SM 10200 H (mod.) / E31780	Phytoplankton chlorophyll-a corrected, uncorrected, and phaeophytin by spectrophotometry	2117237, 2117238, 2117239, 2117240, 2117241
SM 9223 B Quanti-Tray / E84589	E. coli by Colilert-18/QT by AEL in Tampa	2117244, 2117245, 2117246, 2117247, 2117248
SOP-PCR-1.0 / E31780	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	2117242, 2117243
SOP-PCR-1.4 / E31780	Detection and quantification of Ruminant specific Bacteroidetes BacR genetic marker with quantitative polymerase chain reaction	2117242

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
SM 10200 H (mod.)	09/10/2019	09/10/2019 14:18	Yuliya Danyuk	09/11/2019 10:10	Peterson Janvier	2117237, 2117238, 2117239, 2117240
	09/10/2019	09/10/2019 14:18	Yuliya Danyuk	09/11/2019 15:05	Peterson Janvier	2117241
SM 9223 B Quanti-Tray	09/09/2019	09/09/2019 12:48		09/10/2019 08:06	AEL - Tampa	2117244, 2117245, 2117246, 2117247, 2117248
SOP-PCR-1.0	09/10/2019	09/10/2019 14:00	Angela Smith	09/24/2019 08:33	Mailin Lopez	2117242, 2117243
SOP-PCR-1.4	09/10/2019	09/10/2019 14:00	Angela Smith	09/24/2019 09:07	Mailin Lopez	2117242