

Biological Analysis Report

SW-DIST-2019-02-19-03

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 - Tampa
9610 Princess Palm Avenue
Tampa, Florida 33619
DOH Accreditation E84589

Event Description: **SMP - English, Poley Mustang**
Request ID: **RQ-2019-02-18-23**
Customer: **SW-DIST**
Project ID: **SMP**

Send Reports to:
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Certified by: Cheryl A. Swanson, Program Administrator

Date Certified: 26-MAR-2019 15:14



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: Mustang Ranch Creek @ Hopewell

Collection Date/Time: 02/18/2019 11:35

Field ID: G2SW0026

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2063491	SOP-PCR-1.0	HF183-qPCR**	940		GEU/100 mL	P359134	
	SOP-PCR-1.4	BacR-qPCR**	1.45E+05		TSC/100 mL	P359134	MS
2063494	EPA 1603	Escherichia coli-Membrane Filter**	2600		cfu/100 mL		

Sample Location: Poley @ Shepard Rd

Collection Date/Time: 02/18/2019 11:10

Field ID: G2SW0014

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2063492	SOP-PCR-1.0	HF183-qPCR**	280	U	GEU/100 mL	P359134	
	SOP-PCR-1.4	BacR-qPCR**	3.0E+03	U	TSC/100 mL	P359134	MS

Sample Location: English Creek above SR 60

Collection Date/Time: 02/18/2019 12:45

Field ID: G2SW0006

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2063493	SOP-PCR-1.0	HF183-qPCR**	140	U	GEU/100 mL	P359134	
	SOP-PCR-1.4	BacR-qPCR**	5.8E+03	I	TSC/100 mL	P359134	MS
2063496	EPA 1603	Escherichia coli-Membrane Filter**	144		cfu/100 mL		

Sample Location: English Creek above SR 60

Collection Date/Time: 02/18/2019 12:45

Field ID: G2SW0006

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2063490	SM 10200 H (mod.)	Chlorophyll-a, Corrected	0.83	I	ug/L	P359794	
		Phaeophytin-a	0.60	U	ug/L	P359794	
		Chlorophyll-a, Uncorrected	0.98	I	ug/L	P359794	

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: Poley Creek @ SR60

Collection Date/Time: 02/18/2019 11:10

Field ID: G2SW0049

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2068269	EPA 1603	Escherichia coli-Membrane Filter**	550		cfu/100 mL		

Quality Assurance Report Method Blank Results

Reference Method: SM 10200 H (mod.)

Batch ID: P359794

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

Quality Assurance Report Method Blank Results

Reference Method: SOP-PCR-1.0
Batch ID: P359134

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

Reference Method: SOP-PCR-1.4
Batch ID: P359134

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: P359794

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	101		P	84 - 116

Reference Method: SOP-PCR-1.0
Batch ID: P359134

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	126	143	P/P	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P359134

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
BacR-qPCR	103	77.5	P/P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
Batch ID: P359134

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2061397	HF183-qPCR	78.2		P	50 - 175
2061398	HF183-qPCR	92.6		P	50 - 175
2062017	HF183-qPCR	99.3		P	50 - 175
2063491	HF183-qPCR	129		P	50 - 175
2063492	HF183-qPCR	120		P	50 - 175
2063493	HF183-qPCR	124	115	P/P	50 - 175
2063752	HF183-qPCR	95.8		P	50 - 175
2063753	HF183-qPCR	101		P	50 - 175
2063756	HF183-qPCR	68.5		P	50 - 175
2063757	HF183-qPCR	91.0		P	50 - 175
2063758	HF183-qPCR	74.4		P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.4
Batch ID: P359134

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2063491	BacR-qPCR	66.3		P	50 - 175
2063492	BacR-qPCR	81.3		P	50 - 175
2063493	BacR-qPCR	75.4	76.8	P/P	50 - 175
2063752	BacR-qPCR	58.3		P	50 - 175
2063753	BacR-qPCR	59.1		P	50 - 175
2063756	BacR-qPCR	41.4		F	50 - 175
2063757	BacR-qPCR	67.6		P	50 - 175
2063758	BacR-qPCR	44.2		F	50 - 175

Quality Assurance Report Precision

Reference Method: SM 10200 H (mod.)
Batch ID: P359794

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2063790	Chlorophyll-a, Corrected	3.38	Sample	P	0 - 25
2063790	Chlorophyll-a, Uncorrected	5.14	Sample	P	0 - 25

Reference Method: SOP-PCR-1.0
Batch ID: P359134

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2063493	HF183-qPCR	7.25	Spike	P	0 - 25

Reference Method: SOP-PCR-1.4
Batch ID: P359134

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2063493	BacR-qPCR	1.76	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: 2063491
Field Sample ID: G2SW0026

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	105	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	98.2	P	50 - 300

Lab Sample ID: 2063492
Field Sample ID: G2SW0014

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

Lab Sample ID: 2063492
Field Sample ID: G2SW0014

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	128	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	122	P	50 - 300

Lab Sample ID: 2063493
Field Sample ID: G2SW0006

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	114	P	50 - 300
SOP-PCR-1.0	SKETA-qPCR	113	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	109	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	101						3.38	
	Chlorophyll-a, Uncorrected							5.14	
SOP-PCR-1.0	HF183-qPCR	143	126	129	120	124			7.25
				115					
SOP-PCR-1.4	BacR-qPCR	77.5	103	66.3	81.3	75.4			1.76
				76.8					

Reference Method Descriptions

Method / DoH Cert #	Description	Associated Samples
EPA 1603 / E84589	Escherichia coli by membrane filter	2063494, 2063496, 2068269
SM 10200 H (mod.) / E31780	Phytoplankton chlorophyll-a corrected, uncorrected, and phaeophytin by spectrophotometry	2063490
SOP-PCR-1.0 / E31780	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	2063491, 2063492, 2063493
SOP-PCR-1.4 / E31780	Detection and quantification of Ruminant specific Bacteroidetes BacR genetic marker with quantitative polymerase chain reaction	2063491, 2063492, 2063493

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
EPA 1603	02/18/2019	02/18/2019 14:26		02/19/2019 13:32	AEL - Tampa	2063494, 2063496, 2068269
SM 10200 H (mod.)	02/19/2019	02/19/2019 13:45	Jennifer Mihalic	02/28/2019 14:50	Carina A. Tull	2063490
SOP-PCR-1.0	02/19/2019	02/19/2019 14:30	Angela Smith	03/01/2019 11:25	Mailin Lopez	2063491, 2063492, 2063493
SOP-PCR-1.4	02/19/2019	02/19/2019 14:30	Angela Smith	03/04/2019 08:42	Mailin Lopez	2063491, 2063492, 2063493