

Biological Analysis Report

CEN-DIST-2019-08-16-01

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, Inc.
380 North Lake Blvd, Ste 1048
Altamonte Springs, FL 32701
DOH Accreditation E53076

Event Description: **Econ Trib / Roberts**
Request ID: **RQ-2019-07-22-36**
Customer: **CEN-DIST**
Project ID: **SMP**

Send Reports to:
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Certified by: Jacqueline Savage, Environmental Administrator

Date Certified: 30-SEP-2019 11:58

A handwritten signature in black ink that reads "Jackie Savage". The signature is written in a cursive, flowing style.

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: Roberts Branch @ Curryville Rd

Collection Date/Time: 08/15/2019 10:50

Field ID: G3CE0011

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2112427	EPA 1603	Escherichia coli Membrane Filter	520		cfu/100mL		
2112428	SM 10200 H (mod.)	Chlorophyll-a, Corrected	1.6	U	ug/L	P369812	
		Phaeophytin-a	1.8	U	ug/L	P369812	
		Chlorophyll-a, Uncorrected	1.2	U	ug/L	P369812	
2112429	SOP-PCR-1.0	HF183-qPCR**	280	U	GEU/100 mL	P369163	
	SOP-PCR-1.4	BacR-qPCR**	1.2E+04	I	TSC/100 mL	P369163	

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: P369812

Component	Result	Code	Units
Chlorophyll-a, Corrected	0.96	U	ug/L
Chlorophyll-a, Uncorrected	0.70	U	ug/L
Phaeophytin-a	1.0	U	ug/L

Reference Method: SOP-PCR-1.0

Batch ID: P369163

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

Reference Method: SOP-PCR-1.4

Batch ID: P369163

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: P369812

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	85.6		P	84 - 116

Reference Method: SOP-PCR-1.0

Batch ID: P369163

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	133	150	P/P	50 - 175

Reference Method: SOP-PCR-1.4

Batch ID: P369163

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
BacR-qPCR	89.9	72.7	P/P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
Batch ID: P369163

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2112421	HF183-qPCR	119	128	P/P	50 - 175
2112429	HF183-qPCR	109		P	50 - 175
2112732	HF183-qPCR	107		P	50 - 175
2112733	HF183-qPCR	94.1		P	50 - 175
2112870	HF183-qPCR	96.2		P	50 - 175
2113066	HF183-qPCR	109		P	50 - 175
2113067	HF183-qPCR	113		P	50 - 175
2113068	HF183-qPCR	112		P	50 - 175
2113227	HF183-qPCR	106		P	50 - 175
2113228	HF183-qPCR	103		P	50 - 175
2113229	HF183-qPCR	98.0		P	50 - 175
2113559	HF183-qPCR	87.7		P	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P369163

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2112421	BacR-qPCR	79.6	80.8	P/P	50 - 175
2112429	BacR-qPCR	53.9		P	50 - 175
2113066	BacR-qPCR	64.9		P	50 - 175
2113067	BacR-qPCR	61.5		P	50 - 175
2113068	BacR-qPCR	60.5		P	50 - 175

Quality Assurance Report Precision

Reference Method: SM 10200 H (mod.)
Batch ID: P369812

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2112172	Chlorophyll-a, Uncorrected	3.73	Sample	P	0 - 25

Reference Method: SOP-PCR-1.0
Batch ID: P369163

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2112421	HF183-qPCR	7.40	Spike	P	0 - 25

Reference Method: SOP-PCR-1.4
Batch ID: P369163

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2112421	BacR-qPCR	1.52	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: 2112429
Field Sample ID: G3CE0011

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	95.4	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	102	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	85.6							
	Chlorophyll-a, Uncorrected							3.73	
SOP-PCR-1.0	HF183-qPCR	150	133	119	128	109			7.40
				107					
SOP-PCR-1.4	BacR-qPCR	72.7	89.9	79.6	80.8	53.9			1.52
				64.9					

Reference Method Descriptions

Method / DoH Cert #	Description	Associated Samples
EPA 1603 / E53076	E. coli by MF by AEL in Altamonte Springs	2112427
SM 10200 H (mod.) / E31780	Phytoplankton chlorophyll-a corrected, uncorrected, and phaeophytin by spectrophotometry	2112428
SOP-PCR-1.0 / E31780	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	2112429
SOP-PCR-1.4 / E31780	Detection and quantification of Ruminant specific Bacteroidetes BacR genetic marker with quantitative polymerase chain reaction	2112429

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
EPA 1603	08/15/2019	08/15/2019 15:35		08/16/2019 16:00	Advanced Environmental Laboratory	2112427
SM 10200 H (mod.)	08/16/2019	08/16/2019 14:26	Jennifer Mihalic	08/26/2019 15:11	Yuliya Danyuk	2112428
SOP-PCR-1.0	08/16/2019	08/16/2019 13:20	Angela Smith	09/12/2019 14:03	Mailin Lopez	2112429
SOP-PCR-1.4	08/16/2019	08/16/2019 13:20	Angela Smith	09/12/2019 12:51	Mailin Lopez	2112429