

Biological Analysis Report

SW-DIST-2019-07-30-01

Florida Department of Environmental Protection
Central Laboratory
2600 Blair Stone Road
Tallahassee, FL 32399-2400
DOH Accreditation E31780

Overflow Analyses Performed By:
Benchmark Environmental, Inc.
1711 12th Street East
Palmetto, FL 34221
DOH Accreditation E84167

Event Description: **SMP: Run 8**
Request ID: **RQ-2019-07-29-01**
Customer: **SW-DIST**
Project ID: **SMP**

Send Reports to:

FL Dept. of Environmental Protection
FL Dept. of Environmental Protection
13051 N. Telecom Parkway
Temple Terrace, FL 33637-0926
Attn: Ben Paswater

For additional information please contact

Cheryl Swanson - Administrator
Jacqueline Savage - Microbiology
Diane Power - Bench Biology
Andrew Tintle - Aquatic Toxicology
Puja Jasrotia, Ph.D. - Molecular Biology
Amelia Rankin - Taxonomy (Invertebrate & Algal)
Thekkekalathil Chandrasekhar, Ph.D, QA Officer
Phone (850) 245-8177

Certified by: Jacqueline Savage, Environmental Administrator

Date Certified: 19-SEP-2019 13:09



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: LIL MANATEE RIVER N.FORK @ CARLTON LAKE RD

Collection Date/Time: 07/29/2019 09:30

Field ID: G2SW0100

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2107028	SOP-PCR-1.0	HF183-qPCR**	140	U	GEU/100 mL	P369893	
	SOP-PCR-1.4	BacR-qPCR**	5.0E+03	I	TSC/100 mL	P369893	
2107032	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	150		MPN/100 mL		

Sample Location: GAMBLE CREEK @ MULHOLLAND RD

Collection Date/Time: 07/29/2019 10:50

Field ID: G2SW0106

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2107030	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	20		MPN/100 mL		

Sample Location: GAMBLE CREEK @ GULF COURSE RD

Collection Date/Time: 07/29/2019 10:30

Field ID: G2SW0107

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2107031	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	109		MPN/100 mL		

Sample Location: LIL MANATEE RIVER @ SAFFOLD RD

Collection Date/Time: 07/29/2019 09:50

Field ID: G2SW0103

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2107027	SM 10200 H (mod.)	Chlorophyll-a, Corrected	0.82	U	ug/L	P368660	
		Phaeophytin-a	0.90	U	ug/L	P368660	
		Chlorophyll-a, Uncorrected	0.60	U	ug/L	P368660	
2107029	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	146		MPN/100 mL		

Quality Assurance Report Method Blank Results

Reference Method: SM 10200 H (mod.)

Batch ID: P368660

Component	Result	Code	Units
Chlorophyll-a, Corrected	0.82	U	ug/L
Chlorophyll-a, Uncorrected	0.60	U	ug/L
Phaeophytin-a	0.90	U	ug/L

Reference Method: SOP-PCR-1.0

Batch ID: P369893

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

Reference Method: SOP-PCR-1.4

Batch ID: P369893

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: P368660

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

Reference Method: SOP-PCR-1.0
Batch ID: P369893

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	143	171	P/P	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P369893

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
BacR-qPCR	90.1	83.3	P/P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
Batch ID: P369893

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2107028	HF183-qPCR	109		P	50 - 175
2115044	HF183-qPCR	132		P	50 - 175
2115733	HF183-qPCR	118		P	50 - 175
2115734	HF183-qPCR	135		P	50 - 175
2115735	HF183-qPCR	118		P	50 - 175
2115736	HF183-qPCR	129	118	P/P	50 - 175
2115970	HF183-qPCR	116		P	50 - 175
2116025	HF183-qPCR	113		P	50 - 175
2116097	HF183-qPCR	131		P	50 - 175
2116098	HF183-qPCR	124		P	50 - 175
2116099	HF183-qPCR	129		P	50 - 175
2116100	HF183-qPCR	124		P	50 - 175
2116129	HF183-qPCR	119		P	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P369893

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2107028	BacR-qPCR	63.2		P	50 - 175
2115733	BacR-qPCR	54.9		P	50 - 175
2115734	BacR-qPCR	67.7		P	50 - 175
2115735	BacR-qPCR	60.1		P	50 - 175
2115736	BacR-qPCR	67.2	76.2	P/P	50 - 175
2115970	BacR-qPCR	55.2		P	50 - 175
2116097	BacR-qPCR	86.9		P	50 - 175
2116098	BacR-qPCR	81.4		P	50 - 175
2116099	BacR-qPCR	86.7		P	50 - 175
2116100	BacR-qPCR	85.1		P	50 - 175
2116129	BacR-qPCR	65.0		P	50 - 175

Quality Assurance Report Precision

Reference Method: SM 10200 H (mod.)
Batch ID: P368660

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2106998	Chlorophyll-a, Corrected	0.340	Sample	P	0 - 25
2106998	Chlorophyll-a, Uncorrected	0.895	Sample	P	0 - 25
2106998	Phaeophytin-a	3.84	Sample	P	0 - 25

Reference Method: SOP-PCR-1.0
Batch ID: P369893

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2115736	HF183-qPCR	8.63	Spike	P	0 - 25

Reference Method: SOP-PCR-1.4
Batch ID: P369893

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2115736	BacR-qPCR	12.6	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: 2107028
Field Sample ID: G2SW0100

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	108	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	108	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	90.2						0.340	
	Chlorophyll-a, Uncorrected							0.895	
	Phaeophytin-a							3.84	
SOP-PCR-1.0	HF183-qPCR	171	143	118	135	118			8.63
SOP-PCR-1.4	BacR-qPCR	83.3	90.1	54.9	67.7	60.1			12.6
				67.2					

Reference Method Descriptions

Method / DoH Cert #	Description	Associated Samples
SM 10200 H (mod.) / E31780	Phytoplankton chlorophyll-a corrected, uncorrected, and phaeophytin by spectrophotometry	2107027

Reference Method Descriptions

Method / DoH Cert #	Description	<u>Associated Samples</u>
SM 9223 B Quanti-Tray / E84167	E. coli by Colilert/QT by BEA in Palmetto	2107029, 2107030, 2107031, 2107032
SOP-PCR-1.0 / E31780	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	2107028
SOP-PCR-1.4 / E31780	Detection and quantification of Ruminant specific Bacteroidetes BacR genetic marker with quantitative polymerase chain reaction	2107028

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
SM 10200 H (mod.)	07/30/2019	07/30/2019 14:00	Peterson Janvier	08/06/2019 10:16	Yuliya Danyuk	2107027
SM 9223 B Quanti-Tray	07/29/2019	07/29/2019 13:26		07/30/2019 14:32	Benchmark EnviroAnalytical - Palmetto	2107029, 2107030, 2107031, 2107032
SOP-PCR-1.0	07/30/2019	07/30/2019 15:30	Angela Smith	09/12/2019 10:16	Mailin Lopez	2107028
SOP-PCR-1.4	07/30/2019	07/30/2019 15:30	Angela Smith	09/13/2019 14:57	Mailin Lopez	2107028