

Biological Analysis Report

SO-DIST-2019-07-12-01

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

Overflow Analyses Performed By:
Benchmark EnviroAnalytical, Inc. - North Port
1001 Corporate Avenue, Suite 102
North Port, FL 34289
DOH Accreditation E85086

Event Description: **P Cricks 2**
Request ID: **RQ-2019-07-08-39**
Customer: **SO-DIST**
Project ID: **SMP**

Send Reports to:
FL Dept. of Environmental Protection
FLDEP South District Office
2295 Victoria Ave. Suite 364
Fort Myers, FL 33901-3381
Attn: Maryann Dugan

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: Puja Jasrotia, Environmental Manager

Date Certified: 08-AUG-2019 14:37



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: HawthorneCreek@Reynolds

Collection Date/Time: 07/11/2019 09:20

Field ID: G3SD0011

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2102567	SM 10200 H (mod.)	Chlorophyll-a, Corrected	1.9	I	ug/L	P367865	
		Phaeophytin-a	1.5	U	ug/L	P367865	
		Chlorophyll-a, Uncorrected	2.9	I	ug/L	P367865	
2102571	SOP-PCR-1.0	HF183-qPCR**	140	U	GEU/100 mL	P367051	
2102575	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray	369		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.

SOP-PCR-1.0: Precision data unavailable for Analysis Batch A92934

Sample Location: Hog Bay @ CR 763

Collection Date/Time: 07/11/2019 10:00

Field ID: G3SD0096

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2102568	SM 10200 H (mod.)	Chlorophyll-a, Corrected	8.3		ug/L	P367865	
		Phaeophytin-a	1.7	I	ug/L	P367865	
		Chlorophyll-a, Uncorrected	9.6		ug/L	P367865	
2102572	SOP-PCR-1.0	HF183-qPCR**	280	U	GEU/100 mL	P367051	
	SOP-PCR-1.4	BacR-qPCR**	6.1E+03	I	TSC/100 mL	P367051	
2102576	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray	119		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.

SOP-PCR-1.0: Precision data unavailable for Analysis Batch A92934

Sample Location: MyrtleSlough@PinelIsland

Collection Date/Time: 07/11/2019 10:45

Field ID: G3SD0095

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2102570	SM 10200 H (mod.)	Chlorophyll-a, Corrected	2.7	I	ug/L	P367865	
		Phaeophytin-a	1.5	U	ug/L	P367865	
		Chlorophyll-a, Uncorrected	2.6	I	ug/L	P367865	
2102574	SOP-PCR-4.0	HF183 - Human Source Marker**	Below Threshold			P366861	
2102578	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray	121		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: Myrtle Slough @ SR 31

Collection Date/Time: 07/11/2019 11:40

Field ID: G3SD0097

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2102569	SM 10200 H (mod.)	Chlorophyll-a, Corrected	1.5	I	ug/L	P367865	
		Phaeophytin-a	0.75	U	ug/L	P367865	
		Chlorophyll-a, Uncorrected	1.9		ug/L	P367865	
2102573	SOP-PCR-1.0	HF183-qPCR**	140	U	GEU/100 mL	P367051	
	SOP-PCR-1.4	BacR-qPCR**	6.32E+04		TSC/100 mL	P367051	
2102577	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray	282		MPN/100 mL		

Field ID: G3SD0097

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
-----------	-------------	-----------	--------	------	-------	----------	----------

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.
 SOP-PCR-1.0: Precision data unavailable for Analysis Batch A92934

Quality Assurance Report Method Blank Results

Reference Method: SM 10200 H (mod.)
 Batch ID: P367865

Component	Result	Code	Units
Chlorophyll-a, Corrected	0.69	U	ug/L
Chlorophyll-a, Uncorrected	0.50	U	ug/L
Phaeophytin-a	0.75	U	ug/L

Reference Method: SOP-PCR-1.0
 Batch ID: P367051

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

Reference Method: SOP-PCR-1.4
 Batch ID: P367051

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: P367865

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	85.9		P	84 - 116

Reference Method: SOP-PCR-1.0
 Batch ID: P367051

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	146	157	P/P	50 - 175

Reference Method: SOP-PCR-1.4
 Batch ID: P367051

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
BacR-qPCR	93.4	69.5	P/P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
 Batch ID: P367051

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2102571	HF183-qPCR	109		P	50 - 175
2102572	HF183-qPCR	122		P	50 - 175
2102573	HF183-qPCR	133		P	50 - 175

Reference Method: SOP-PCR-1.4
 Batch ID: P367051

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2102572	BacR-qPCR	74.4		P	50 - 175
2102573	BacR-qPCR	90.7		P	50 - 175
2102670	BacR-qPCR	78.6		P	50 - 175
2102671	BacR-qPCR	77.9	70.7	P/P	50 - 175

Quality Assurance Report Precision

Reference Method: SM 10200 H (mod.)
 Batch ID: P367865

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2102630	Chlorophyll-a, Uncorrected	1.49	Sample	P	0 - 25
2102630	Phaeophytin-a	10.9	Sample	P	0 - 25

Reference Method: SOP-PCR-1.4
 Batch ID: P367051

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2102671	BacR-qPCR	9.60	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: 2102571
 Field Sample ID: G3SD0011

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	98.8	P	50 - 300

Lab Sample ID: 2102572
 Field Sample ID: G3SD0096

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	93.5	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	98.5	P	50 - 300

Quality Assurance Report Surrogates

Lab Sample ID: 2102573
Field Sample ID: G3SD0097

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	109	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	107	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	85.9							
	Chlorophyll-a, Uncorrected							1.49	
	Phaeophytin-a							10.9	
SOP-PCR-1.0	HF183-qPCR	157	146	109	122	133			
SOP-PCR-1.4	BacR-qPCR	69.5	93.4	74.4	90.7	78.6			9.60
				77.9					

Reference Method Descriptions

Method / DoH Cert #	Description	Associated Samples
SM 10200 H (mod.) / E31780	Phytoplankton chlorophyll-a corrected, uncorrected, and phaeophytin by spectrophotometry	2102567, 2102568, 2102569, 2102570
SM 9223 Quanti-Tray / E85086	E. coli by Colilert/QT by BEA South in North Port	2102575, 2102576, 2102577, 2102578
SOP-PCR-1.0 / E31780	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	2102571, 2102572, 2102573
SOP-PCR-1.4 / E31780	Detection and quantification of Ruminant specific Bacteroidetes BacR genetic marker with quantitative polymerase chain reaction	2102572, 2102573
SOP-PCR-4.0 / E31780	Preparation of Samples by Filtration and held for qPCR Analysis	2102574

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
SM 10200 H (mod.)	07/12/2019	07/12/2019 13:55	Edwin Gonzalez-Soto	07/23/2019 15:55	Peterson Janvier	2102567, 2102568, 2102569, 2102570
SM 9223 Quanti-Tray	07/11/2019	07/11/2019 14:27		07/12/2019 15:08	Benchmark EnviroAnalytical in North Port	2102575, 2102576, 2102577, 2102578
SOP-PCR-1.0	07/12/2019	07/12/2019 14:00	Angela Smith	07/22/2019 10:15	Mailin Lopez	2102571, 2102572, 2102573
SOP-PCR-1.4	07/12/2019	07/12/2019 14:00	Angela Smith	07/26/2019 08:30	Mailin Lopez	2102572, 2102573
SOP-PCR-4.0	07/12/2019	07/12/2019 14:00	Angela Smith	07/23/2019 11:29	Sarah Menz	2102574