

Biological Analysis Report

SO-DIST-2019-07-10-01

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 1**
Request ID: **RQ-2019-07-01-48**
Customer: **SO-DIST**
Project ID: **SMP**

Send Reports to:
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FLDEP South District Office
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Fort Myers, FL 33901-3381
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Certified by: Puja Jasrotia, Environmental Manager

Date Certified: 07-AUG-2019 14:32



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: Shell Cr@HathawayPark

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

Field ID: G3SD0120

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2101333	SM 10200 H (mod.)	Chlorophyll-a, Corrected	17	A	ug/L	P367675	
		Phaeophytin-a	4.0	I	ug/L	P367675	
		Chlorophyll-a, Uncorrected	20	A	ug/L	P367675	
2101341	SOP-PCR-1.0	HF183-qPCR**	140	U	GEU/100 mL	P366693	
	SOP-PCR-1.4	BacR-qPCR**	1.5E+03	U	TSC/100 mL	P366693	
2101344	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray	41		MPN/100 mL		

Ref. Method and Comment:

SM 10200 H (mod.): Laboratory duplicate result is 3.7 I for Phaeophytin-a. Precision data is unavailable due to the small amount of analyte in the QC sample.

SOP-PCR-1.4: Precision data unavailable for Analysis Batch A92844

Sample Location: PrairieCreek@WshgtnLoop

Collection Date/Time: 07/09/2019 10:15

Field ID: G3SD0067

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2101334	SM 10200 H (mod.)	Chlorophyll-a, Corrected	1.6	U	ug/L	P367675	
		Phaeophytin-a	1.8	U	ug/L	P367675	
		Chlorophyll-a, Uncorrected	1.2	I	ug/L	P367675	
2101342	SOP-PCR-1.0	HF183-qPCR**	140	U	GEU/100 mL	P366693	
	SOP-PCR-1.4	BacR-qPCR**	3.6E+03	I	TSC/100 mL	P366693	
2101345	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray	135		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-1.4: Precision data unavailable for Analysis Batch A92844

Sample Location: RunofftoPeaceRiverSite3

Collection Date/Time: 07/09/2019 11:05

Field ID: G3SD0132

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2101336	SM 10200 H (mod.)	Chlorophyll-a, Corrected	1.6	U	ug/L	P367675	
		Phaeophytin-a	1.8	U	ug/L	P367675	
		Chlorophyll-a, Uncorrected	1.5	I	ug/L	P367675	
2101343	SOP-PCR-1.0	HF183-qPCR**	1.80E+03		GEU/100 mL	P366693	
	SOP-PCR-1.4	BacR-qPCR**	1.5E+03	U	TSC/100 mL	P366693	
2101347	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray	537		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-1.4: Precision data unavailable for Analysis Batch A92844

Sample Location: PrairieCreek@WshgtnLoop

Collection Date/Time: 07/09/2019 11:45

Field ID: G3SD0067_DUP

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2101335	SM 10200 H (mod.)	Chlorophyll-a, Corrected	1.7	I	ug/L	P367675	
		Phaeophytin-a	1.8	U	ug/L	P367675	
		Chlorophyll-a, Uncorrected	2.0	I	ug/L	P367675	

Field ID: G3SD0067_DUP

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2101346	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray	110		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: BobcatCreekPreserveEast

Collection Date/Time: 07/09/2019 12:35

Field ID: G3SD0131

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2101337	SM 10200 H (mod.)	Chlorophyll-a, Corrected	1.2	U	ug/L	P367676	
		Phaeophytin-a	1.4	U	ug/L	P367676	
		Chlorophyll-a, Uncorrected	1.4	I	ug/L	P367676	
2101348	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray	602		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: P367675

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: SM 10200 H (mod.)

Batch ID: P367676

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: P366693

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

Reference Method: SOP-PCR-1.4

Batch ID: P366693

Component	Result	Code	Units
BacR-qPCR	3.0E+03	U	TSC/100 mL

Quality Assurance Report Laboratory Control Sample Accuracy

Quality Assurance Report Laboratory Control Sample Accuracy

Reference Method: SM 10200 H (mod.)
Batch ID: P367675

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	87.5		P	84 - 116

Reference Method: SM 10200 H (mod.)
Batch ID: P367676

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	87.9		P	84 - 116

Reference Method: SOP-PCR-1.0
Batch ID: P366693

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	131	174	P/P	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P366693

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
BacR-qPCR	96.9	71.0	P/P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
Batch ID: P366693

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2098288	HF183-qPCR	105		P	50 - 175
2098810	HF183-qPCR	119		P	50 - 175
2098811	HF183-qPCR	115		P	50 - 175
2098812	HF183-qPCR	116		P	50 - 175
2098958	HF183-qPCR	102	97.5	P/P	50 - 175
2100818	HF183-qPCR	88.3		P	50 - 175
2100819	HF183-qPCR	117		P	50 - 175
2101175	HF183-qPCR	118		P	50 - 175
2101341	HF183-qPCR	107		P	50 - 175
2101342	HF183-qPCR	95.8		P	50 - 175
2101343	HF183-qPCR	101		P	50 - 175
2101411	HF183-qPCR	99.2		P	50 - 175
2101593	HF183-qPCR	79.5		P	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P366693

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2098288	BacR-qPCR	80.2		P	50 - 175
2098810	BacR-qPCR	80.4		P	50 - 175
2098811	BacR-qPCR	91.9		P	50 - 175
2098812	BacR-qPCR	89.9		P	50 - 175
2100818	BacR-qPCR	65.3		P	50 - 175
2101175	BacR-qPCR	85.9		P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.4
Batch ID: P366693

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2101341	BacR-qPCR	86.2		P	50 - 175
2101342	BacR-qPCR	70.6		P	50 - 175
2101343	BacR-qPCR	75.1		P	50 - 175
2101593	BacR-qPCR	68.9		P	50 - 175

Quality Assurance Report Precision

Reference Method: SM 10200 H (mod.)
Batch ID: P367675

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2101333	Chlorophyll-a, Corrected	3.70	Sample	P	0 - 25
2101333	Chlorophyll-a, Uncorrected	4.36	Sample	P	0 - 25

Reference Method: SM 10200 H (mod.)
Batch ID: P367676

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2101401	Chlorophyll-a, Corrected	7.52	Sample	P	0 - 25
2101401	Chlorophyll-a, Uncorrected	6.65	Sample	P	0 - 25

Reference Method: SOP-PCR-1.0
Batch ID: P366693

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2098958	HF183-qPCR	4.53	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: 2101341
Field Sample ID: G3SD0120

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	98.5	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	103	P	50 - 300

Lab Sample ID: 2101342
Field Sample ID: G3SD0067

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	92.7	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	98.8	P	50 - 300

Quality Assurance Report Surrogates

Lab Sample ID: 2101343
Field Sample ID: G3SD0132

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	101	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	103	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	87.5						3.70	
	Chlorophyll-a, Corrected	87.9						7.52	
	Chlorophyll-a, Uncorrected							4.36	
	Chlorophyll-a, Uncorrected							6.65	
SOP-PCR-1.0	HF183-qPCR	174	131	119	115	116			4.53
SOP-PCR-1.4	BacR-qPCR	71.0	96.9	80.4	91.9	89.9			
				65.3					

Reference Method Descriptions

Method / DoH Cert #	Description	Associated Samples
SM 10200 H (mod.) / E31780	Phytoplankton chlorophyll-a corrected, uncorrected, and phaeophytin by spectrophotometry	2101333, 2101334, 2101335, 2101336, 2101337
SM 9223 Quanti-Tray / E85086	E. coli by Colilert/QT by BEA South in North Port	2101344, 2101345, 2101346, 2101347, 2101348
SOP-PCR-1.0 / E31780	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	2101341, 2101342, 2101343
SOP-PCR-1.4 / E31780	Detection and quantification of Ruminant specific Bacteroidetes BacR genetic marker with quantitative polymerase chain reaction	2101341, 2101342, 2101343

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
SM 10200 H (mod.)	07/10/2019	07/10/2019 13:01	Yuliya Danyuk	07/18/2019 11:15	Yuliya Danyuk	2101333, 2101334, 2101335, 2101336
	07/10/2019	07/10/2019 13:01	Yuliya Danyuk	07/19/2019 09:30	Peterson Janvier	2101337
SM 9223 Quanti-Tray	07/09/2019	07/09/2019 15:36		07/10/2019 15:53	Benchmark EnviroAnalytical in North Port	2101344, 2101345, 2101346, 2101347, 2101348
SOP-PCR-1.0	07/10/2019	07/10/2019 15:20	Sarah Menz	07/17/2019 08:29	Mailin Lopez	2101341, 2101342, 2101343
SOP-PCR-1.4	07/10/2019	07/10/2019 15:20	Sarah Menz	07/17/2019 14:49	Mailin Lopez	2101341, 2101342, 2101343