

Biological Analysis Report

SO-DIST-2020-06-26-04

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

Overflow Analyses Performed By:
Sanders Laboratories
10090 Bavaria Road
Ft. Myers, FL 33913
DOH Accreditation E85457

Event Description: **2020 SMP : Peace River**

Request ID: **RQ-2020-04-13-15**

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: Chris Hormuth

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Certified by: Puja Jasrotia, Environmental Manager

Date Certified: 23-JUL-2020 13:55



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: BeeBranch@LoblollyBayRd

Collection Date/Time: 06/25/2020 10:15

Field ID: G3SD0085

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2179073	SM 10200 H (mod.)	Chlorophyll-a, Corrected	2.1	U	ug/L	P384068	
		Phaeophytin-a	2.2	U	ug/L	P384068	
		Chlorophyll-a, Uncorrected	1.5	U	ug/L	P384068	
2179074	SOP-PCR-1.0	HF183-qPCR**	280	U	GEU/100 mL	P383562	
	SOP-PCR-1.4	BacR-qPCR**	7.93E+04		TSC/100 mL	P383562	
	SOP-PCR-1.5	DG3-qPCR**	2.6E+03	U	TSC/100 mL	P383562	
2179075	Colilert-18	Escherichia Coli-Quanti-Tray	186		MPN/100 mL		

Quality Assurance Report Method Blank Results

Reference Method: SM 10200 H (mod.)
 Batch ID: P384068

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: P383562

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

Reference Method: SOP-PCR-1.4
 Batch ID: P383562

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

Reference Method: SOP-PCR-1.5
 Batch ID: P383562

Component	Result	Code	Units
DG3-qPCR	2.6E+03	U	TSC/100 mL

Quality Assurance Report Laboratory Control Sample Accuracy

Reference Method: SM 10200 H (mod.)
 Batch ID: P384068

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	103		P	84 - 116

Reference Method: SOP-PCR-1.0
 Batch ID: P383562

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	107	121	P/P	50 - 175

Quality Assurance Report Laboratory Control Sample Accuracy

Reference Method: SOP-PCR-1.4
Batch ID: P383562

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
BacR-qPCR	98.0	95.9	P/P	50 - 175

Reference Method: SOP-PCR-1.5
Batch ID: P383562

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
DG3-qPCR	98.1	107	P/P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
Batch ID: P383562

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2177136	HF183-qPCR	104	103	P/P	50 - 175
2178000	HF183-qPCR	94.6		P	50 - 175
2178238	HF183-qPCR	89.6		P	50 - 175
2178439	HF183-qPCR	119		P	50 - 175
2178479	HF183-qPCR	101		P	50 - 175
2179074	HF183-qPCR	75.6		P	50 - 175
2179545	HF183-qPCR	84.4		P	50 - 175
2179853	HF183-qPCR	98.2		P	50 - 175
2180812	HF183-qPCR	85.0		P	50 - 175
2180813	HF183-qPCR	90.3		P	50 - 175
2180814	HF183-qPCR	116		P	50 - 175
2180887	HF183-qPCR	101		P	50 - 175
2180888	HF183-qPCR	106		P	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P383562

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2177136	BacR-qPCR	69.9	73.1	P/P	50 - 175
2178000	BacR-qPCR	67.4		P	50 - 175
2178238	BacR-qPCR	96.0		P	50 - 175
2178439	BacR-qPCR	87.5		P	50 - 175
2178479	BacR-qPCR	84.2		P	50 - 175
2179074	BacR-qPCR	56.1		P	50 - 175
2179545	BacR-qPCR	60.2		P	50 - 175
2179853	BacR-qPCR	77.4		P	50 - 175
2180812	BacR-qPCR	71.9		P	50 - 175
2180813	BacR-qPCR	63.9		P	50 - 175
2180814	BacR-qPCR	84.2		P	50 - 175
2180887	BacR-qPCR	74.0		P	50 - 175
2180888	BacR-qPCR	70.0		P	50 - 175

Reference Method: SOP-PCR-1.5
Batch ID: P383562

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
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Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.5
Batch ID: P383562

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2177136	DG3-qPCR	90.4	96.1	P/P	50 - 175
2178000	DG3-qPCR	86.0		P	50 - 175
2178238	DG3-qPCR	91.5		P	50 - 175
2178439	DG3-qPCR	100		P	50 - 175
2178479	DG3-qPCR	96.9		P	50 - 175
2179074	DG3-qPCR	80.3		P	50 - 175
2179545	DG3-qPCR	79.8		P	50 - 175
2179853	DG3-qPCR	99.2		P	50 - 175
2180812	DG3-qPCR	89.8		P	50 - 175
2180813	DG3-qPCR	81.1		P	50 - 175
2180814	DG3-qPCR	104		P	50 - 175
2180887	DG3-qPCR	94.5		P	50 - 175
2180888	DG3-qPCR	97.4		P	50 - 175

Quality Assurance Report Precision

Reference Method: SM 10200 H (mod.)
Batch ID: P384068

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2179330	Chlorophyll-a, Corrected	4.33	Sample	P	0 - 25
2179330	Chlorophyll-a, Uncorrected	3.87	Sample	P	0 - 25
2179330	Phaeophytin-a	0.177	Sample	P	0 - 25

Reference Method: SOP-PCR-1.0
Batch ID: P383562

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2177136	HF183-qPCR	0.785	Spike	P	0 - 25

Reference Method: SOP-PCR-1.4
Batch ID: P383562

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2177136	BacR-qPCR	4.51	Spike	P	0 - 25

Reference Method: SOP-PCR-1.5
Batch ID: P383562

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2177136	DG3-qPCR	6.21	Spike	P	0 - 25

* Sample, spike and/or laboratory control sample precision (LCS) is reported.

Quality Assurance Report Surrogates

Lab Sample ID: 2179074
Field Sample ID: G3SD0085

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	56.4	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	64.0	P	50 - 300
SOP-PCR-1.5	SKETA-qPCR	53.6	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	103						4.33	
	Chlorophyll-a, Uncorrected							3.87	
	Phaeophytin-a							0.177	
SOP-PCR-1.0	HF183-qPCR	121	107	104	103	94.6			0.785
				89.6					
SOP-PCR-1.4	BacR-qPCR	95.9	98.0	69.9	73.1	67.4			4.51
				96.0					
SOP-PCR-1.5	DG3-qPCR	107	98.1	90.4	96.1	86.0			6.21
				91.5					

Reference Method Descriptions

Method	Description	Associated Samples
Colilert-18	E. coli by Colilert-18/QT by Sanders in Fort Myers	2179075
SM 10200 H (mod.)	Phytoplankton chlorophyll-a corrected, uncorrected, and phaeophytin by spectrophotometry	2179073
SOP-PCR-1.0	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	2179074
SOP-PCR-1.4	Detection and quantification of Ruminant specific Bacteroidetes BacR genetic marker with quantitative polymerase chain reaction	2179074
SOP-PCR-1.5	Detection and quantification of canine-specific DG3 Bacteroides genetic marker with quantitative polymerase chain reaction	2179074

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
Colilert-18	06/25/2020	06/25/2020 15:50		06/26/2020 09:57	Sanders-Fort Myers	2179075
SM 10200 H (mod.)	06/26/2020	06/26/2020 14:55	Carina A. Tull	07/02/2020 09:20	Jessica Fetgatter	2179073
SOP-PCR-1.0	06/26/2020	06/26/2020 15:25	Angela Smith	07/14/2020 14:24	Lauren Campbell	2179074
SOP-PCR-1.4	06/26/2020	06/26/2020 15:25	Angela Smith	07/15/2020 10:42	Lauren Campbell	2179074
SOP-PCR-1.5	06/26/2020	06/26/2020 15:25	Angela Smith	07/15/2020 12:16	Lauren Campbell	2179074