

Biological Analysis Report

SO-DIST-2023-04-07-01

Florida DEP 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: **2023 SMP : Land O' Lakes**
Request ID: **RQ-2023-03-27-31**
Customer: **SO-DIST**
Project ID: **SMP**

Send Reports to:
FL Dept. of Environmental Protection
FLDEP South District Office
2295 Victoria Ave. Suite 314
Fort Myers, FL 33901-3381
Attn: William Mullen

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Certified by: Sarah Menz, Environmental Administrator

Date Certified: 05-MAY-2023 12:36

A handwritten signature in dark ink, appearing to be 'Sarah Menz', with a long, sweeping horizontal stroke extending to the right.

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: Bonnet Creek at SR 64

Collection Date/Time: 04/06/2023 10:00

Field ID: G4SD0107

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2399287	SM 10150 B	Chlorophyll-a, Corrected	3.6	I	ug/L	P428810	
		Phaeophytin-a	1.8	U	ug/L	P428810	
		Chlorophyll-a, Uncorrected	4.4		ug/L	P428810	
2399288	SOP-PCR-1.0	HF183-qPCR**	3.3E+03	U	TSC/100 mL	P427755	
	SOP-PCR-1.4	BacR-qPCR**	1.75E+05		TSC/100 mL	P427755	
	SOP-PCR-1.5	DG3-qPCR**	2.6E+03	U	TSC/100 mL	P427755	
2399291	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray	3448		MPN/100 mL		

Ref. Method and Comment:

SM 10150 B: 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: Red Water Lake at Mid

Collection Date/Time: 04/06/2023 12:10

Field ID: G4SD0224

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2399285	SM 10150 B	Chlorophyll-a, Corrected	9.9		ug/L	P428810	
		Phaeophytin-a	1.8	U	ug/L	P428810	
		Chlorophyll-a, Uncorrected	10		ug/L	P428810	
2399289	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray	10		MPN/100 mL		

Ref. Method and Comment:

SM 10150 B: 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: Lake June in Winter

Collection Date/Time: 04/06/2023 12:55

Field ID: G4SD0186

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2399283	SM 10150 B	Chlorophyll-a, Corrected	5.7		ug/L	P428810	
		Phaeophytin-a	0.90	U	ug/L	P428810	
		Chlorophyll-a, Uncorrected	6.1		ug/L	P428810	

Ref. Method and Comment:

SM 10150 B: 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: LAKEPLACIDOUTLETATSR70W

Collection Date/Time: 04/06/2023 14:05

Field ID: G4SD0213

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2399286	SM 10150 B	Chlorophyll-a, Corrected	24		ug/L	P428810	
		Phaeophytin-a	12		ug/L	P428810	
		Chlorophyll-a, Uncorrected	32		ug/L	P428810	
2399290	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray	7270		MPN/100 mL		

Ref. Method and Comment:

SM 10150 B: 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: Lake Glenada

Collection Date/Time: 04/06/2023 10:35

Field ID: G4SD0179

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2399284	SM 10150 B	Chlorophyll-a, Corrected	45		ug/L	P428810	
		Phaeophytin-a	4.9	U	ug/L	P428810	
		Chlorophyll-a, Uncorrected	46		ug/L	P428810	

Ref. Method and Comment:
SM 10150 B: 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 10150 B
Batch ID: P428810

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: P427755

Component	Result	Code	Units
HF183-qPCR	3.3E+03	U	TSC/100 mL

Reference Method: SOP-PCR-1.4
Batch ID: P427755

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

Reference Method: SOP-PCR-1.5
Batch ID: P427755

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

Quality Assurance Report
Laboratory Control Sample Accuracy

Reference Method: SM 10150 B
Batch ID: P428810

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	92.3		P	84 - 116

Reference Method: SOP-PCR-1.0
Batch ID: P427755

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	89.1	100	P/P	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P427755

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

Reference Method: SOP-PCR-1.4
Batch ID: P427755

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
BacR-qPCR	96.5	87.5	P/P	50 - 175

Reference Method: SOP-PCR-1.5
Batch ID: P427755

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
DG3-qPCR	102	95.5	P/P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
Batch ID: P427755

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2396582	HF183-qPCR	76.2	83.8	P/P	50 - 175
2396696	HF183-qPCR	79.9		P	50 - 175
2397523	HF183-qPCR	91.2		P	50 - 175
2397524	HF183-qPCR	95.7		P	50 - 175
2397894	HF183-qPCR	81.3		P	50 - 175
2398313	HF183-qPCR	79.7		P	50 - 175
2398618	HF183-qPCR	84.6		P	50 - 175
2398619	HF183-qPCR	85.8		P	50 - 175
2398964	HF183-qPCR	85.9		P	50 - 175
2398965	HF183-qPCR	85.9		P	50 - 175
2399288	HF183-qPCR	80.6		P	50 - 175
2399389	HF183-qPCR	85.9		P	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P427755

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2396582	BacR-qPCR	78.1	79.7	P/P	50 - 175
2396696	BacR-qPCR	87.9		P	50 - 175
2397523	BacR-qPCR	94.2		P	50 - 175
2397524	BacR-qPCR	88.4		P	50 - 175
2397894	BacR-qPCR	80.9		P	50 - 175
2398313	BacR-qPCR	80.2		P	50 - 175
2398618	BacR-qPCR	84.2		P	50 - 175
2398619	BacR-qPCR	83.2		P	50 - 175
2398964	BacR-qPCR	79.2		P	50 - 175
2398965	BacR-qPCR	82.8		P	50 - 175
2399288	BacR-qPCR	66.8		P	50 - 175
2399389	BacR-qPCR	81.0		P	50 - 175

Reference Method: SOP-PCR-1.5
Batch ID: P427755

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2396582	DG3-qPCR	92.7	88.0	P/P	50 - 175
2396696	DG3-qPCR	94.2		P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.5
Batch ID: P427755

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2397523	DG3-qPCR	105		P	50 - 175
2397524	DG3-qPCR	102		P	50 - 175
2397894	DG3-qPCR	96.7		P	50 - 175
2398313	DG3-qPCR	95.7		P	50 - 175
2398618	DG3-qPCR	97.2		P	50 - 175
2398619	DG3-qPCR	98.7		P	50 - 175
2398964	DG3-qPCR	86.6		P	50 - 175
2398965	DG3-qPCR	95.9		P	50 - 175
2399288	DG3-qPCR	90.7		P	50 - 175
2399389	DG3-qPCR	108		P	50 - 175

Quality Assurance Report Precision

Reference Method: SM 10150 B
Batch ID: P428810

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2399479	Chlorophyll-a, Corrected	4.38	Sample	P	0 - 25
2399479	Chlorophyll-a, Uncorrected	4.88	Sample	P	0 - 25

Reference Method: SOP-PCR-1.0
Batch ID: P427755

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2396582	HF183-qPCR	9.55	Spike	P	0 - 25

Reference Method: SOP-PCR-1.4
Batch ID: P427755

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2396582	BacR-qPCR	1.99	Spike	P	0 - 25

Reference Method: SOP-PCR-1.5
Batch ID: P427755

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2396582	DG3-qPCR	5.17	Spike	P	0 - 25

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

Quality Assurance Report Surrogates

Quality Assurance Report Surrogates

Lab Sample ID: 2399288
 Field Sample ID: G4SD0107

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	97.4	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	117	P	50 - 300
SOP-PCR-1.5	SKETA-qPCR	99.7	P	50 - 300

Quality Assurance Report Summary

Ref. Method	Analyte	LCS % Recovery		MS % Recovery			Precision SMP	MS
SM 10150 B	Chlorophyll-a, Corrected	92.3					4.38	
	Chlorophyll-a, Uncorrected						4.88	
SOP-PCR-1.0	HF183-qPCR	100	89.1	76.2	83.8	79.9		9.55
SOP-PCR-1.4	BacR-qPCR	87.5	96.5	78.1	79.7	87.9		1.99
SOP-PCR-1.5	DG3-qPCR	95.5	102	92.7	88.0	94.2		5.17
				105				

Reference Method Descriptions

Method	Description	Associated Samples
SM 10150 B	Phytoplankton chlorophyll-a corrected, uncorrected, and phaeophytin by spectrophotometry	2399283, 2399284, 2399285, 2399286, 2399287
SM 9223 Quanti-Tray	E. coli by Colilert/QT by BEA South in North Port	2399289, 2399290, 2399291
SOP-PCR-1.0	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	2399288
SOP-PCR-1.4	Detection and quantification of Ruminant specific Bacteroidetes BacR genetic marker with quantitative polymerase chain reaction	2399288
SOP-PCR-1.5	Detection and quantification of canine-specific DG3 Bacteroides genetic marker with quantitative polymerase chain reaction	2399288

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
SM 10150 B	04/07/2023	04/07/2023 14:33	Tynan A Branch	04/18/2023 09:41	Callie Williams	2399283, 2399284, 2399285, 2399286, 2399287
SM 9223 Quanti-Tray	04/06/2023	04/06/2023 16:01		04/07/2023 16:16	Benchmark EnviroAnalytical in North Port	2399289, 2399290, 2399291
SOP-PCR-1.0	04/07/2023	04/07/2023 15:35	Claudia Castro	04/14/2023 10:06	Roberta Tatti	2399288
SOP-PCR-1.4	04/07/2023	04/07/2023 15:35	Claudia Castro	04/14/2023 15:28	Roberta Tatti	2399288
SOP-PCR-1.5	04/07/2023	04/07/2023 15:35	Claudia Castro	04/17/2023 11:02	Roberta Tatti	2399288