

Biological Analysis Report

SO-DIST-2023-02-09-01

Florida DEP Laboratory
2600 Blair Stone Road
Tallahassee, FL 32399-2400
DOH Accreditation E31780

Event Description: **2023 SMP : Land O' Lakes**
Request ID: **RQ-2023-01-30-52**
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

For additional information please contact
Cheryl Swanson - Administrator
Sarah Menz, Ph.D. - Bench Biology & Aquatic
Toxicity
Puja Jasrotia, Ph.D. - Molecular Biology & Taxonomy
Thekkekalathil Chandrasekhar, Ph.D, QA Officer
Phone (850) 245-8177

Certified by: Puja Jasrotia, Environmental Administrator

Date Certified: 16-MAR-2023 13:24



NON-CONFORMANCE REPORT INCLUDED

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 and Molecular.

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 @ SR 64

Collection Date/Time: 02/07/2023 10:15

Field ID: G4SD0107

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2386142	SOP-PCR-1.0	HF183-qPCR**	1.1E+03	UQ	TSC/100 mL	P425465	
	SOP-PCR-1.4	BacR-qPCR**	1.43E+04	Q	TSC/100 mL	P425465	
	SOP-PCR-1.5	DG3-qPCR**	860	UQ	TSC/100 mL	P425465	

Ref. Method and Comment:

SOP-PCR-1.0: See non-conformance report # 9744

SOP-PCR-1.4: See non-conformance report # 9744

SOP-PCR-1.5: See non-conformance report # 9744.

Sample Location: Lake Placid Outlet at SR 70 W

Collection Date/Time: 02/07/2023 13:35

Field ID: G4SD0213

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2386141	SM 10200 H (mod.)	Chlorophyll-a, Corrected	1.3	I	ug/L	P425742	
		Phaeophytin-a	0.90	U	ug/L	P425742	
		Chlorophyll-a, Uncorrected	1.8	I	ug/L	P425742	RPD

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.

Non-Conformance Report

NCR ID: 9744

Event(s)

SO-DIST-2023-02-09-01

Job(s)

TLH-2023-02-09-11

Sample(s)

2386142

Test(s)

NCR Type: SHIPPING/RECEIVING

NCR Category: Samples arrived late

Observation: Samples arrived late, samples received within the acceptable temperature range.

Resolution: Samples will be qualified as appropriate.
Biology and Chemistry supervisors were notified by email.

Authorized by/Date: Kyle Klug 2/16/2023

The Non-Conformance Report details exceptions or problems encountered with the events/jobs/samples/test.

Please address questions to:

Chemistry Colin Wright (850) 245-8085
Biology Cheryl Swanson (850) 245-8177

Quality Assurance Report

Method Blank Results

Reference Method: SM 10200 H (mod.)

Batch ID: P425742

Component	Result	Code	Units
Chlorophyll-a, Corrected	0.89	U	ug/L
Chlorophyll-a, Uncorrected	0.65	U	ug/L
Phaeophytin-a	0.98	U	ug/L

Quality Assurance Report Method Blank Results

Reference Method: SOP-PCR-1.0
Batch ID: P425465

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

Reference Method: SOP-PCR-1.4
Batch ID: P425465

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

Reference Method: SOP-PCR-1.5
Batch ID: P425465

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: P425742

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	96.2		P	84 - 116

Reference Method: SOP-PCR-1.0
Batch ID: P425465

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	97.0	96.4	P/P	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P425465

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
BacR-qPCR	105	111	P/P	50 - 175

Reference Method: SOP-PCR-1.5
Batch ID: P425465

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
DG3-qPCR	99.0	108	P/P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
Batch ID: P425465

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2385071	HF183-qPCR	83.0	85.7	P/P	50 - 175
2385072	HF183-qPCR	88.6		P	50 - 175
2385073	HF183-qPCR	93.8		P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
Batch ID: P425465

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2385804	HF183-qPCR	78.9		P	50 - 175
2385851	HF183-qPCR	76.8		P	50 - 175
2385907	HF183-qPCR	79.0		P	50 - 175
2386142	HF183-qPCR	73.6		P	50 - 175
2386201	HF183-qPCR	94.3		P	50 - 175
2386202	HF183-qPCR	83.9		P	50 - 175
2386211	HF183-qPCR	105		P	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P425465

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2385071	BacR-qPCR	94.3	95.2	P/P	50 - 175
2385072	BacR-qPCR	93.6		P	50 - 175
2385073	BacR-qPCR	92.9		P	50 - 175
2385804	BacR-qPCR	88.0		P	50 - 175
2385851	BacR-qPCR	71.4		P	50 - 175
2385907	BacR-qPCR	81.5		P	50 - 175
2386142	BacR-qPCR	70.2		P	50 - 175
2386201	BacR-qPCR	89.3		P	50 - 175
2386202	BacR-qPCR	80.3		P	50 - 175
2386211	BacR-qPCR	102	75.9	P/P	50 - 175

Reference Method: SOP-PCR-1.5
Batch ID: P425465

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2385071	DG3-qPCR	89.9	96.2	P/P	50 - 175
2385072	DG3-qPCR	100		P	50 - 175
2385073	DG3-qPCR	101		P	50 - 175
2385428	DG3-qPCR	103		P	50 - 175
2385429	DG3-qPCR	105		P	50 - 175
2385430	DG3-qPCR	99.3		P	50 - 175
2385804	DG3-qPCR	104		P	50 - 175
2385851	DG3-qPCR	96.8		P	50 - 175
2385907	DG3-qPCR	104		P	50 - 175
2386142	DG3-qPCR	92.1		P	50 - 175
2386201	DG3-qPCR	105		P	50 - 175
2386202	DG3-qPCR	88.9		P	50 - 175
2386211	DG3-qPCR	105		P	50 - 175

Quality Assurance Report Precision

Reference Method: SM 10200 H (mod.)
Batch ID: P425742

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2385871	Chlorophyll-a, Uncorrected	37.4	Sample	F	0 - 25

Quality Assurance Report Precision

Reference Method: SOP-PCR-1.0
Batch ID: P425465

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2385071	HF183-qPCR	3.22	Spike	P	0 - 25

Reference Method: SOP-PCR-1.4
Batch ID: P425465

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2385071	BacR-qPCR	1.00	Spike	P	0 - 25

Reference Method: SOP-PCR-1.5
Batch ID: P425465

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2385071	DG3-qPCR	6.79	Spike	P	0 - 25

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

Quality Assurance Report Surrogates

Lab Sample ID: 2386142
Field Sample ID: G4SD0107

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	75.4	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	90.9	P	50 - 300
SOP-PCR-1.5	SKETA-qPCR	72.8	P	50 - 300

Quality Assurance Report Summary

Ref. Method	Analyte	LCS % Recovery		MS % Recovery			Precision SMP	MS
SM 10200 H (mod.)	Chlorophyll-a, Corrected	96.2					37.4	
	Chlorophyll-a, Uncorrected							
SOP-PCR-1.0	HF183-qPCR	96.4	97.0	83.0	85.7	88.6		3.22
				93.8				
SOP-PCR-1.4	BacR-qPCR	111	105	75.9	80.3	102		1.00
				94.3				
SOP-PCR-1.5	DG3-qPCR	108	99.0	89.9	96.2	100		6.79
				101				

Reference Method Descriptions

Method	Description	Associated Samples
SM 10200 H (mod.)	Phytoplankton chlorophyll-a corrected, uncorrected, and phaeophytin by spectrophotometry	2386141

Reference Method Descriptions

Method	Description	<u>Associated Samples</u>
SOP-PCR-1.0	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	2386142
SOP-PCR-1.4	Detection and quantification of Ruminant specific Bacteroidetes BacR genetic marker with quantitative polymerase chain reaction	2386142
SOP-PCR-1.5	Detection and quantification of canine-specific DG3 Bacteroides genetic marker with quantitative polymerase chain reaction	2386142

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
SM 10200 H (mod.)	02/09/2023	02/09/2023 10:59	Callie Williams	02/13/2023 09:17	Aidan Cronin	2386141
SOP-PCR-1.0	02/09/2023	02/09/2023 11:05	Claudia Castro	02/20/2023 23:05	Roberta Tatti	2386142
SOP-PCR-1.4	02/09/2023	02/09/2023 11:05	Claudia Castro	02/20/2023 12:18	Roberta Tatti	2386142
SOP-PCR-1.5	02/09/2023	02/09/2023 11:05	Claudia Castro	02/20/2023 15:32	Roberta Tatti	2386142