

Biological Analysis Report

SE-DIST-2022-12-08-01

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

Event Description: **Lake Clarke and Blank**
Request ID: **RQ-2022-11-14-34**
Customer: **SE-DIST**
Project ID: **SMP**

Send Reports to:
FL Dept. of Environmental Protection
FL Dept. of Environmental Protection
2199 South Rock Road
Fort Pierce, FL 34945
Attn: Jordan Skaggs

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: Sarah Menz, Environmental Administrator

Date Certified: 06-JAN-2023 12:46

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, Microbiology 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: Lake Clarke East of Wagner Circle

Collection Date/Time: 12/07/2022 11:15

Field ID: G3SE0066

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2373801	SM 10200 H (mod.)	Chlorophyll-a, Corrected	11		ug/L	P423599	
		Phaeophytin-a	1.8	I	ug/L	P423599	
		Chlorophyll-a, Uncorrected	13		ug/L	P423599	
2373803	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	17.3	Q	MPN/100 mL	P423587	
2373805	SOP-PCR-1.0	HF183-qPCR**	830	U	TSC/100 mL	P423278	
	SOP-PCR-1.2	GULL2-qPCR**	900	U	TSC/100 mL	P423278	
	SOP-PCR-1.3	GFD-qPCR**	180	U	TSC/100 mL	P423278	
	SOP-PCR-1.4	BacR-qPCR**	760	U	TSC/100 mL	P423278	
	SOP-PCR-1.5	DG3-qPCR**	640	U	TSC/100 mL	P423278	

Ref. Method and Comment:

SM 9223 B Quanti-Tray: Precision data is unavailable for E. coli due to the small amount of analyte in the QC sample for the batch.

Sample Location: Lake Clarke East of Edgewater Circle

Collection Date/Time: 12/07/2022 11:30

Field ID: G3SE0065

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2373802	SM 10200 H (mod.)	Chlorophyll-a, Corrected	4.9		ug/L	P423690	
		Phaeophytin-a	2.2	I	ug/L	P423690	
		Chlorophyll-a, Uncorrected	6.4		ug/L	P423690	RPD
2373804	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	20.3	Q	MPN/100 mL	P423587	
2373806	SOP-PCR-1.0	HF183-qPCR**	1.7E+03	U	TSC/100 mL	P423278	
	SOP-PCR-1.2	GULL2-qPCR**	1.8E+03	U	TSC/100 mL	P423278	
	SOP-PCR-1.3	GFD-qPCR**	710	I	TSC/100 mL	P423278	
	SOP-PCR-1.4	BacR-qPCR**	1.5E+03	U	TSC/100 mL	P423278	
	SOP-PCR-1.5	DG3-qPCR**	1.3E+03	U	TSC/100 mL	P423278	

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.

SM 9223 B Quanti-Tray: Precision data is unavailable for E. coli due to the small amount of analyte in the QC sample for the batch.

SOP-PCR-1.3: Duplicate was analyzed and result is 360 U.

Quality Assurance Report Method Blank Results

Reference Method: SM 10200 H (mod.)

Batch ID: P423599

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: P423690

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

Quality Assurance Report Method Blank Results

Reference Method: SM 9223 B Quanti-Tray
Batch ID: P423587

Component	Result	Code	Units
Escherichia Coli-Quanti-Tray	1.0	U	MPN/100 mL

Reference Method: SOP-PCR-1.0
Batch ID: P423278

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

Reference Method: SOP-PCR-1.2
Batch ID: P423278

Component	Result	Code	Units
GULL2-qPCR	3.6E+03	U	TSC/100 mL

Reference Method: SOP-PCR-1.3
Batch ID: P423278

Component	Result	Code	Units
GFD-qPCR	720	U	TSC/100 mL

Reference Method: SOP-PCR-1.4
Batch ID: P423278

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

Reference Method: SOP-PCR-1.5
Batch ID: P423278

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: P423599

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	98.9		P	84 - 116

Reference Method: SM 10200 H (mod.)
Batch ID: P423690

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	96.9		P	84 - 116

Reference Method: SOP-PCR-1.0
Batch ID: P423278

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

Reference Method: SOP-PCR-1.0
Batch ID: P423278

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	102	90.2	P/P	50 - 175

Reference Method: SOP-PCR-1.2
Batch ID: P423278

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
GULL2-qPCR	95.8	99.3	P/P	50 - 300

Reference Method: SOP-PCR-1.3
Batch ID: P423278

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
GFD-qPCR	87.6	93.1	P/P	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P423278

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
BacR-qPCR	130	84.5	P/P	50 - 175

Reference Method: SOP-PCR-1.5
Batch ID: P423278

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
DG3-qPCR	87.6	88.0	P/P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
Batch ID: P423278

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2373805	HF183-qPCR	91.3	88.9	P	50 - 175
2373806	HF183-qPCR	85.8		P/P	50 - 175
2374161	HF183-qPCR	85.6	88.9	P	50 - 175
2374542	HF183-qPCR	85.4		P	50 - 175
2374653	HF183-qPCR	88.0	88.9	P	50 - 175
2374657	HF183-qPCR	95.7		P	50 - 175
2374926	HF183-qPCR	87.9	88.9	P	50 - 175
2374927	HF183-qPCR	87.7		P	50 - 175
2374941	HF183-qPCR	85.4	88.9	P	50 - 175
2375190	HF183-qPCR	71.0		P	50 - 175
2375636	HF183-qPCR	83.6	88.9	P	50 - 175

Reference Method: SOP-PCR-1.2
Batch ID: P423278

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2373805	GULL2-qPCR	77.1	91.3	P	50 - 300
2373806	GULL2-qPCR	89.8		P/P	50 - 300

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.2

Batch ID: P423278

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2374653	GULL2-qPCR	74.7		P	50 - 300
2374657	GULL2-qPCR	88.9		P	50 - 300
2374926	GULL2-qPCR	93.1		P	50 - 300
2374927	GULL2-qPCR	90.6		P	50 - 300
2374941	GULL2-qPCR	96.7		P	50 - 300
2375190	GULL2-qPCR	56.4		P	50 - 300

Reference Method: SOP-PCR-1.3

Batch ID: P423278

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2373805	GFD-qPCR	93.3		P	50 - 175
2373806	GFD-qPCR	94.2	113	P/P	50 - 175
2374653	GFD-qPCR	84.8		P	50 - 175
2374657	GFD-qPCR	91.5		P	50 - 175
2374926	GFD-qPCR	91.4		P	50 - 175
2374927	GFD-qPCR	92.2		P	50 - 175
2374941	GFD-qPCR	102		P	50 - 175
2375190	GFD-qPCR	76.6		P	50 - 175

Reference Method: SOP-PCR-1.4

Batch ID: P423278

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2373805	BacR-qPCR	104		P	50 - 175
2373806	BacR-qPCR	99.8	102	P/P	50 - 175
2374161	BacR-qPCR	96.6		P	50 - 175
2374542	BacR-qPCR	85.9		P	50 - 175
2374653	BacR-qPCR	83.0		P	50 - 175
2374657	BacR-qPCR	95.7		P	50 - 175
2374926	BacR-qPCR	93.5		P	50 - 175
2374927	BacR-qPCR	96.4		P	50 - 175
2374941	BacR-qPCR	90.4		P	50 - 175
2375190	BacR-qPCR	72.2		P	50 - 175
2375636	BacR-qPCR	77.2		P	50 - 175

Reference Method: SOP-PCR-1.5

Batch ID: P423278

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2373805	DG3-qPCR	89.8		P	50 - 175
2373806	DG3-qPCR	103	99.6	P/P	50 - 175
2374161	DG3-qPCR	87.6		P	50 - 175
2374542	DG3-qPCR	90.5		P	50 - 175
2374653	DG3-qPCR	89.5		P	50 - 175
2374657	DG3-qPCR	94.4		P	50 - 175
2374926	DG3-qPCR	92.8		P	50 - 175
2374927	DG3-qPCR	92.1		P	50 - 175
2374941	DG3-qPCR	92.4		P	50 - 175
2375190	DG3-qPCR	91.4		P	50 - 175
2375636	DG3-qPCR	83.0		P	50 - 175

Quality Assurance Report Precision

Reference Method: SM 10200 H (mod.)
 Batch ID: P423599

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2373568	Chlorophyll-a, Corrected	4.62	Sample	P	0 - 25
2373568	Chlorophyll-a, Uncorrected	2.09	Sample	P	0 - 25
2373568	Phaeophytin-a	16.3	Sample	P	0 - 25

Reference Method: SM 10200 H (mod.)
 Batch ID: P423690

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2373847	Chlorophyll-a, Uncorrected	26.5	Sample	F	0 - 25

Reference Method: SOP-PCR-1.0
 Batch ID: P423278

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2373806	HF183-qPCR	3.56	Spike	P	0 - 25

Reference Method: SOP-PCR-1.2
 Batch ID: P423278

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2373806	GULL2-qPCR	1.65	Spike	P	0 - 25

Reference Method: SOP-PCR-1.3
 Batch ID: P423278

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2373806	GFD-qPCR	18.2	Spike	P	0 - 25

Reference Method: SOP-PCR-1.4
 Batch ID: P423278

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2373806	BacR-qPCR	2.46	Spike	P	0 - 25

Reference Method: SOP-PCR-1.5
 Batch ID: P423278

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2373806	DG3-qPCR	3.44	Spike	P	0 - 25

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

Quality Assurance Report Surrogates

Lab Sample ID: 2373805
Field Sample ID: G3SE0066

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	98.6	P	50 - 300
SOP-PCR-1.2	SKETA-qPCR	109	P	50 - 300
SOP-PCR-1.3	SKETA-qPCR	105	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	107	P	50 - 300
SOP-PCR-1.5	SKETA-qPCR	96.9	P	50 - 300

Lab Sample ID: 2373806
Field Sample ID: G3SE0065

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	99.5	P	50 - 300
SOP-PCR-1.0	SKETA-qPCR	94.7	P	50 - 300
SOP-PCR-1.2	SKETA-qPCR	109	P	50 - 300
SOP-PCR-1.2	SKETA-qPCR	105	P	50 - 300
SOP-PCR-1.3	SKETA-qPCR	108	P	50 - 300
SOP-PCR-1.3	SKETA-qPCR	111	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	114	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	109	P	50 - 300
SOP-PCR-1.5	SKETA-qPCR	105	P	50 - 300
SOP-PCR-1.5	SKETA-qPCR	103	P	50 - 300

Quality Assurance Report Summary

Ref. Method	Analyte	LCS % Recovery			MS % Recovery			LCS	Precision	MS
									SMP	
SM 10200 H (mod.)	Chlorophyll-a, Corrected	98.9							4.62	
	Chlorophyll-a, Corrected	96.9								
	Chlorophyll-a, Uncorrected								2.09	
	Chlorophyll-a, Uncorrected								26.5	
	Phaeophytin-a								16.3	
SOP-PCR-1.0	HF183-qPCR	90.2	102		91.3	85.8	88.9			3.56
SOP-PCR-1.2	GULL2-qPCR	99.3	95.8		85.6					
					77.1	89.8	91.3			1.65
SOP-PCR-1.3	GFD-qPCR	93.1	87.6		74.7					
					93.3	94.2	113			18.2
SOP-PCR-1.4	BacR-qPCR	84.5	130		84.8					
					104	99.8	102			2.46
SOP-PCR-1.5	DG3-qPCR	88.0	87.6		96.6					
					89.8	103	99.6			3.44
					87.6					

Reference Method Descriptions

Method	Description	Associated Samples
SM 10200 H (mod.)	Phytoplankton chlorophyll-a corrected, uncorrected, and phaeophytin by spectrophotometry	2373801, 2373802
SM 9223 B Quanti-Tray	Escherichia coli by Colilert 18 Quanti-Tray method - non-chlorinated water systems	2373803, 2373804
SOP-PCR-1.0	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	2373805, 2373806
SOP-PCR-1.2	Detection and quantification of coastal bird specific Catellicoccus marimammalium Gull2 genetic marker with quantitative polymerase chain reaction	2373805, 2373806

Reference Method Descriptions

Method	Description	<u>Associated Samples</u>
SOP-PCR-1.3	Detection and quantification of bird specific Helicobacter GFD genetic marker with quantitative polymerase chain reaction	2373805, 2373806
SOP-PCR-1.4	Detection and quantification of Ruminant specific Bacteroidetes BacR genetic marker with quantitative polymerase chain reaction	2373805, 2373806
SOP-PCR-1.5	Detection and quantification of canine-specific DG3 Bacteroides genetic marker with quantitative polymerase chain reaction	2373805, 2373806

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
SM 10200 H (mod.)	12/08/2022	12/08/2022 14:10	Aidan Cronin	12/13/2022 15:43	Megan Higbee	2373801
	12/08/2022	12/08/2022 14:10	Aidan Cronin	12/16/2022 09:03	Megan Higbee	2373802
SM 9223 B Quanti-Tray	12/08/2022	12/08/2022 16:40	Tynan A Branch	12/09/2022 11:01	Callie Williams	2373803, 2373804
SOP-PCR-1.0	12/08/2022	12/08/2022 13:40	Claudia Castro	12/19/2022 14:57	DeAsia Armster	2373805, 2373806
SOP-PCR-1.2	12/08/2022	12/08/2022 13:40	Claudia Castro	12/21/2022 08:50	Roberta Tatti	2373805, 2373806
SOP-PCR-1.3	12/08/2022	12/08/2022 13:40	Claudia Castro	12/21/2022 12:32	Roberta Tatti	2373805, 2373806
SOP-PCR-1.4	12/08/2022	12/08/2022 13:40	Claudia Castro	12/20/2022 09:17	Roberta Tatti	2373805, 2373806
SOP-PCR-1.5	12/08/2022	12/08/2022 13:40	Claudia Castro	12/20/2022 12:47	DeAsia Armster	2373805, 2373806