

Biological Analysis Report

CEN-DIST-2023-09-13-02

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

Event Description: **Run 9**
Request ID: **RQ-2023-09-11-33**
Customer: **CEN-DIST**
Project ID: **SMP**

Send Reports to:
FL Dept. of Environmental Protection
FL Dept. of Environmental Protection
3319 Maguire Blvd., Suite 232
Orlando, FL 32803-3767
Attn: Matthew Bearden

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Certified by: Puja Jasrotia, Environmental Administrator

Date Certified: 26-OCT-2023 11:13

A handwritten signature in black ink, appearing to be 'Puja Jasrotia', with a horizontal line underneath it.

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: MATTRESS DRAIN AT ROCKRIDGE RD

Collection Date/Time: 09/12/2023 10:31

Field ID: G4CE0247

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2436360	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	115.3	Q	MPN/100 mL	P435149	
2436362	SOP-PCR-1.0	HF183-qPCR**	1.7E+03	U	TSC/100 mL	P434608	
	SOP-PCR-1.4	BacR-qPCR**	4.5E+03	I	TSC/100 mL	P434608	
	SOP-PCR-1.5	DG3-qPCR**	1.3E+03	U	TSC/100 mL	P434608	LCS

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: BIG CREEK AT LAKE NELLIE RD

Collection Date/Time: 09/12/2023 11:36

Field ID: G1CE0117

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2436358	SM 10150 B	Chlorophyll-a, Corrected	2.4	U	ug/L	P435104	
		Phaeophytin-a	2.6	U	ug/L	P435104	
		Chlorophyll-a, Uncorrected	1.8	U	ug/L	P435104	
2436359	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	148.3	Q	MPN/100 mL	P435149	

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.
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: DAVENPORT CREEK AT HIGHWAY 545

Collection Date/Time: 09/12/2023 13:06

Field ID: G4CE0023

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2436361	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	1732.9	Q	MPN/100 mL	P435149	

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.

Quality Assurance Report

Method Blank Results

Reference Method: SM 10150 B
Batch ID: P435104

Component	Result	Code	Units
Chlorophyll-a, Corrected	0.96	U	ug/L
Chlorophyll-a, Uncorrected	0.70	U	ug/L
Phaeophytin-a	1.0	U	ug/L

Reference Method: SM 9223 B Quanti-Tray
Batch ID: P435149

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

Reference Method: SOP-PCR-1.0
Batch ID: P434608

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

Quality Assurance Report Method Blank Results

Reference Method: SOP-PCR-1.4
Batch ID: P434608

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

Reference Method: SOP-PCR-1.5
Batch ID: P434608

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: P435104

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	90.1		P	84 - 116

Reference Method: SOP-PCR-1.0
Batch ID: P434608

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	87.0	76.4	P/P	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P434608

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
BacR-qPCR	108	69.2	P/P	50 - 175

Reference Method: SOP-PCR-1.5
Batch ID: P434608

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
DG3-qPCR	48.1	66.1	F/P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
Batch ID: P434608

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2434239	HF183-qPCR	74.6	79.8	P/P	50 - 175
2435446	HF183-qPCR	81.5		P	50 - 175
2435447	HF183-qPCR	97.5		P	50 - 175
2435448	HF183-qPCR	85.0		P	50 - 175
2435679	HF183-qPCR	83.4		P	50 - 175
2435726	HF183-qPCR	104		P	50 - 175
2435727	HF183-qPCR	92.2		P	50 - 175
2435728	HF183-qPCR	93.7		P	50 - 175
2435729	HF183-qPCR	95.1		P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
Batch ID: P434608

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2435730	HF183-qPCR	101		P	50 - 175
2435731	HF183-qPCR	88.0		P	50 - 175
2436362	HF183-qPCR	81.4		P	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P434608

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2434239	BacR-qPCR	62.8	58.5	P/P	50 - 175
2435446	BacR-qPCR	90.9		P	50 - 175
2435447	BacR-qPCR	89.9		P	50 - 175
2435448	BacR-qPCR	87.0		P	50 - 175
2435679	BacR-qPCR	59.8		P	50 - 175
2436362	BacR-qPCR	65.1		P	50 - 175

Reference Method: SOP-PCR-1.5
Batch ID: P434608

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2434239	DG3-qPCR	83.2	80.3	P/P	50 - 175
2435446	DG3-qPCR	91.1		P	50 - 175
2435447	DG3-qPCR	87.9		P	50 - 175
2435448	DG3-qPCR	91.4		P	50 - 175
2435679	DG3-qPCR	86.1		P	50 - 175
2435726	DG3-qPCR	105		P	50 - 175
2435727	DG3-qPCR	106		P	50 - 175
2435728	DG3-qPCR	100		P	50 - 175
2435729	DG3-qPCR	99.6		P	50 - 175
2435730	DG3-qPCR	104		P	50 - 175
2435731	DG3-qPCR	92.5		P	50 - 175
2436362	DG3-qPCR	86.1		P	50 - 175

Quality Assurance Report Precision

Reference Method: SM 10150 B
Batch ID: P435104

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2436364	Chlorophyll-a, Corrected	0.00359	Sample	P	0 - 25
2436364	Chlorophyll-a, Uncorrected	0.950	Sample	P	0 - 25

Reference Method: SOP-PCR-1.0
Batch ID: P434608

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2434239	HF183-qPCR	6.69	Spike	P	0 - 25

Quality Assurance Report

Precision

Reference Method: SOP-PCR-1.4

Batch ID: P434608

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2434239	BacR-qPCR	6.94	Spike	P	0 - 25

Reference Method: SOP-PCR-1.5

Batch ID: P434608

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2434239	DG3-qPCR	3.51	Spike	P	0 - 25

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

Quality Assurance Report

Surrogates

Lab Sample ID: 2436362

Field Sample ID: G4CE0247

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	73.5	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	89.1	P	50 - 300
SOP-PCR-1.5	SKETA-qPCR	72.7	P	50 - 300

Quality Assurance Report

Summary

Ref. Method	Analyte	LCS % Recovery		MS % Recovery			Precision	MS
							SMP	
SM 10150 B	Chlorophyll-a, Corrected	90.1					0.0035	
	Chlorophyll-a, Uncorrected						0.950	
SOP-PCR-1.0	HF183-qPCR	76.4	87.0	101	88.0	81.4		6.69
SOP-PCR-1.4	BacR-qPCR	108	69.2	74.6	58.5	90.9		6.94
				62.8				
SOP-PCR-1.5	DG3-qPCR	66.1	48.1	89.9	80.3	91.1		3.51
				83.2				
				87.9				

Reference Method Descriptions

Method	Description	Associated Samples
SM 10150 B	Phytoplankton chlorophyll-a corrected, uncorrected, and phaeophytin by spectrophotometry	2436358
SM 9223 B Quanti-Tray	Escherichia coli by Colilert 18 Quanti-Tray method - non-chlorinated water systems	2436359, 2436360, 2436361
SOP-PCR-1.0	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	2436362
SOP-PCR-1.4	Detection and quantification of Ruminant specific Bacteroidetes BacR genetic marker with quantitative polymerase chain reaction	2436362
SOP-PCR-1.5	Detection and quantification of canine-specific DG3 Bacteroides genetic marker with quantitative polymerase chain reaction	2436362

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
SM 10150 B	09/13/2023	09/13/2023 13:59	Megan Higbee	09/14/2023 13:17	Keanu Thijm	2436358
SM 9223 B Quanti-Tray	09/13/2023	09/13/2023 16:33	Callie Williams	09/14/2023 11:53	Megan Higbee	2436359, 2436360, 2436361
SOP-PCR-1.0	09/13/2023	09/13/2023 15:20	Claudia Castro	09/22/2023 16:51	Madeline Gruver	2436362
SOP-PCR-1.4	09/13/2023	09/13/2023 15:20	Claudia Castro	09/26/2023 17:29	Madeline Gruver	2436362
SOP-PCR-1.5	09/13/2023	09/13/2023 15:20	Claudia Castro	09/26/2023 18:37	Madeline Gruver	2436362