

Biological Analysis Report

SE-DIST-2023-03-02-01

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

Event Description: **SMP - Vero Beach**
Request ID: **RQ-2023-02-27-36**
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: Andrew Luering

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: 13-APR-2023 12:02



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: 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: Vero North Canal @ 66th Ave

Collection Date/Time: 03/01/2023 09:40

Field ID: G5SE0069

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2391295	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	27.9	Q	MPN/100 mL	P426703	
2391303	SOP-PCR-1.0	HF183-qPCR**	1.1E+03	U	TSC/100 mL	P426355	
	SOP-PCR-1.2	GULL2-qPCR**	1.2E+03	U	TSC/100 mL	P426355	
	SOP-PCR-1.3	GFD-qPCR**	4.1E+03		TSC/100 mL	P426355	
	SOP-PCR-1.4	BacR-qPCR**	1.0E+03	U	TSC/100 mL	P426355	
	SOP-PCR-1.5	DG3-qPCR**	860	U	TSC/100 mL	P426355	

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: North Canal @US1

Collection Date/Time: 03/01/2023 10:10

Field ID: G5SE0043

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2391296	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	172.3	Q	MPN/100 mL	P426703	
2391304	SOP-PCR-1.0	HF183-qPCR**	1.7E+03	U	TSC/100 mL	P426355	
	SOP-PCR-1.2	GULL2-qPCR**	1.8E+03	U	TSC/100 mL	P426355	
	SOP-PCR-1.3	GFD-qPCR**	1.9E+03		TSC/100 mL	P426355	
	SOP-PCR-1.4	BacR-qPCR**	1.5E+03	U	TSC/100 mL	P426355	
	SOP-PCR-1.5	DG3-qPCR**	1.3E+03	U	TSC/100 mL	P426355	

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: MAIN CANAL @ COUNTRY CLUB DR

Collection Date/Time: 03/01/2023 10:30

Field ID: G5SE0113

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2391297	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	146.7	Q	MPN/100 mL	P426703	

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: Main Canal @ Kings Highway

Collection Date/Time: 03/01/2023 10:55

Field ID: G5SE0115

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2391298	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	33.6	JQ	MPN/100 mL	P426703	

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. See Non-Conformance Report 9780.

Sample Location: South Canal @ 43rd Ave

Collection Date/Time: 03/01/2023 11:10

Field ID: G5SE0087

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2391299	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	313.0	Q	MPN/100 mL	P426703	

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: South Canal @ 20th Ave.

Collection Date/Time: 03/01/2023 11:25

Field ID: G5SE0086

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2391300	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	45.9	Q	MPN/100 mL	P426703	

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.

Non-Conformance Report

NCR ID: 9780

Event(s)	Job(s)	Sample(s)	Test(s)
SE-DIST-2023-03-02-01	TLH-2023-03-02-71	2391298	ECOLI-18QT

NCR Type: ANALYSIS NCR Category: Analytical Measurement

Observation: Marine sample was not diluted at least tenfold according to the method.
Resolution: Sample result will be J-qualified and customers, Jordan Skaggs and Andrew Luering, notified.

Authorized by/Date: Cheryl A. Swanson 3/10/2023

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 9223 B Quanti-Tray
Batch ID: P426703

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

Reference Method: SOP-PCR-1.0
Batch ID: P426355

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

Reference Method: SOP-PCR-1.2
Batch ID: P426355

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

Reference Method: SOP-PCR-1.3
Batch ID: P426355

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

Reference Method: SOP-PCR-1.4
Batch ID: P426355

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

Quality Assurance Report
Method Blank Results

Reference Method: SOP-PCR-1.5
Batch ID: P426355

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

Quality Assurance Report
Laboratory Control Sample Accuracy

Reference Method: SOP-PCR-1.0
Batch ID: P426355

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	80.7	89.8	P/P	50 - 175

Reference Method: SOP-PCR-1.2
Batch ID: P426355

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
GULL2-qPCR	103	90.2	P/P	50 - 300

Reference Method: SOP-PCR-1.3
Batch ID: P426355

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
GFD-qPCR	104	97.2	P/P	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P426355

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
BacR-qPCR	101	162	P/P	50 - 175

Reference Method: SOP-PCR-1.5
Batch ID: P426355

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
DG3-qPCR	102	153	P/P	50 - 175

Quality Assurance Report
Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
Batch ID: P426355

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2389955	HF183-qPCR	90.2	90.2	P/P	50 - 175
2390307	HF183-qPCR	86.1		P	50 - 175
2390320	HF183-qPCR	85.9		P	50 - 175
2390519	HF183-qPCR	92.1		P	50 - 175
2390521	HF183-qPCR	91.8		P	50 - 175
2390522	HF183-qPCR	94.4		P	50 - 175
2390646	HF183-qPCR	94.4		P	50 - 175
2390722	HF183-qPCR	90.1		P	50 - 175
2391303	HF183-qPCR	92.3		P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
Batch ID: P426355

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2391304	HF183-qPCR	94.1		P	50 - 175
2391448	HF183-qPCR	94.4		P	50 - 175
2391599	HF183-qPCR	96.8		P	50 - 175

Reference Method: SOP-PCR-1.2
Batch ID: P426355

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2389955	GULL2-qPCR	86.4	89.0	P/P	50 - 300
2390307	GULL2-qPCR	63.5		P	50 - 300
2390320	GULL2-qPCR	81.0		P	50 - 300
2390521	GULL2-qPCR	73.9		P	50 - 300
2390646	GULL2-qPCR	68.9		P	50 - 300
2390722	GULL2-qPCR	86.3		P	50 - 300
2391303	GULL2-qPCR	96.7		P	50 - 300
2391304	GULL2-qPCR	73.0		P	50 - 300

Reference Method: SOP-PCR-1.3
Batch ID: P426355

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2389955	GFD-qPCR	103	110	P/P	50 - 175
2390307	GFD-qPCR	90.6		P	50 - 175
2390320	GFD-qPCR	95.9		P	50 - 175
2390521	GFD-qPCR	94.7		P	50 - 175
2390646	GFD-qPCR	85.4		P	50 - 175
2390722	GFD-qPCR	103		P	50 - 175
2391303	GFD-qPCR	95.7		P	50 - 175
2391304	GFD-qPCR	94.2		P	50 - 175
2391448	GFD-qPCR	99.5		P	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P426355

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2389955	BacR-qPCR	92.9	88.9	P/P	50 - 175
2390307	BacR-qPCR	71.2		P	50 - 175
2390320	BacR-qPCR	78.3		P	50 - 175
2390519	BacR-qPCR	90.8		P	50 - 175
2390521	BacR-qPCR	72.2		P	50 - 175
2390522	BacR-qPCR	80.0		P	50 - 175
2390646	BacR-qPCR	78.6		P	50 - 175
2390722	BacR-qPCR	89.2		P	50 - 175
2391303	BacR-qPCR	92.1		P	50 - 175
2391304	BacR-qPCR	82.3		P	50 - 175
2391448	BacR-qPCR	85.6		P	50 - 175
2391599	BacR-qPCR	91.4		P	50 - 175

Reference Method: SOP-PCR-1.5
Batch ID: P426355

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: P426355

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2389955	DG3-qPCR	97.8	101	P/P	50 - 175
2390307	DG3-qPCR	92.4		P	50 - 175
2390320	DG3-qPCR	97.1		P	50 - 175
2390519	DG3-qPCR	105		P	50 - 175
2390521	DG3-qPCR	103		P	50 - 175
2390522	DG3-qPCR	98.8		P	50 - 175
2390646	DG3-qPCR	97.9		P	50 - 175
2390722	DG3-qPCR	107		P	50 - 175
2391303	DG3-qPCR	106		P	50 - 175
2391304	DG3-qPCR	99.6		P	50 - 175
2391599	DG3-qPCR	103		P	50 - 175

Quality Assurance Report Precision

Reference Method: SOP-PCR-1.0
Batch ID: P426355

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2389955	HF183-qPCR	0.0101	Spike	P	0 - 25

Reference Method: SOP-PCR-1.2
Batch ID: P426355

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2389955	GULL2-qPCR	3.04	Spike	P	0 - 25

Reference Method: SOP-PCR-1.3
Batch ID: P426355

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2389955	GFD-qPCR	6.98	Spike	P	0 - 25

Reference Method: SOP-PCR-1.4
Batch ID: P426355

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2389955	BacR-qPCR	4.32	Spike	P	0 - 25

Reference Method: SOP-PCR-1.5
Batch ID: P426355

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2389955	DG3-qPCR	3.35	Spike	P	0 - 25

Quality Assurance Report Precision

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

Quality Assurance Report Surrogates

Lab Sample ID: 2391303
Field Sample ID: G5SE0069

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	106	P	50 - 300
SOP-PCR-1.2	SKETA-qPCR	90.7	P	50 - 300
SOP-PCR-1.3	SKETA-qPCR	92.3	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	95.3	P	50 - 300
SOP-PCR-1.5	SKETA-qPCR	101	P	50 - 300

Lab Sample ID: 2391304
Field Sample ID: G5SE0043

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	93.9	P	50 - 300
SOP-PCR-1.2	SKETA-qPCR	85.5	P	50 - 300
SOP-PCR-1.3	SKETA-qPCR	83.3	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	90.4	P	50 - 300
SOP-PCR-1.5	SKETA-qPCR	89.4	P	50 - 300

Quality Assurance Report Summary

Ref. Method	Analyte	LCS % Recovery		MS % Recovery			Precision SMP	MS
SOP-PCR-1.0	HF183-qPCR	89.8	80.7	90.2	90.2	86.1		0.0101
SOP-PCR-1.2	GULL2-qPCR	90.2	103	85.9	86.4	89.0		3.04
SOP-PCR-1.3	GFD-qPCR	97.2	104	81.0	103	110		6.98
SOP-PCR-1.4	BacR-qPCR	162	101	95.9	92.9	88.9		4.32
SOP-PCR-1.5	DG3-qPCR	153	102	78.3	97.8	101		3.35
				97.1	92.4			

Reference Method Descriptions

Method	Description	Associated Samples
SM 9223 B Quanti-Tray	Escherichia coli by Colilert 18 Quanti-Tray method - non-chlorinated water systems	2391295, 2391296, 2391297, 2391298, 2391299, 2391300
SOP-PCR-1.0	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	2391303, 2391304
SOP-PCR-1.2	Detection and quantification of coastal bird specific Catellicoccus marimammalium Gull2 genetic marker with quantitative polymerase chain reaction	2391303, 2391304
SOP-PCR-1.3	Detection and quantification of bird specific Helicobacter GFD genetic marker with quantitative polymerase chain reaction	2391303, 2391304
SOP-PCR-1.4	Detection and quantification of Ruminant specific Bacteroidetes BacR genetic marker with quantitative polymerase chain reaction	2391303, 2391304

Reference Method Descriptions

Method	Description	<u>Associated Samples</u>
SOP-PCR-1.5	Detection and quantification of canine-specific DG3 Bacteroides genetic marker with quantitative polymerase chain reaction	2391303, 2391304

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
SM 9223 B Quanti-Tray	03/02/2023	03/02/2023 15:29	Megan Higbee	03/03/2023 11:36	Megan Higbee	2391295, 2391296, 2391297, 2391298, 2391299, 2391300
SOP-PCR-1.0	03/02/2023	03/02/2023 14:00	Claudia Castro	03/20/2023 10:10	Roberta Tatti	2391303, 2391304
SOP-PCR-1.2	03/02/2023	03/02/2023 14:00	Claudia Castro	03/28/2023 14:01	Jennifer Mihalic	2391303, 2391304
SOP-PCR-1.3	03/02/2023	03/02/2023 14:00	Claudia Castro	03/27/2023 15:57	Jennifer Mihalic	2391303, 2391304
SOP-PCR-1.4	03/02/2023	03/02/2023 14:00	Claudia Castro	03/21/2023 09:39	Roberta Tatti	2391303, 2391304
SOP-PCR-1.5	03/02/2023	03/02/2023 14:00	Claudia Castro	03/21/2023 10:50	Roberta Tatti	2391303, 2391304