

Biological Analysis Report

SO-DIST-2023-07-26-01

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: SARASOTA**

Request ID: **RQ-2023-07-24-44**

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: Monica Jaeger

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Certified by: Puja Jasrotia, Environmental Administrator

Date Certified: 07-SEP-2023 09:53



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: 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: NORTH CREEK (TIDAL) AT US 41

Collection Date/Time: 07/25/2023 09:40

Field ID: G3SD0089

Matrix: W-SURF-SLT

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2426803	Enterolert/QT	Enterococci-Quanti-Tray	719		MPN/100 mL		

Sample Location: NORTH CREEK BY OAKS CREEK CT

Collection Date/Time: 07/25/2023 10:10

Field ID: G3SD0064

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2426802	SOP-PCR-1.0	HF183-qPCR**	3.3E+03	U	TSC/100 mL	P432974	
	SOP-PCR-1.2	GULL2-qPCR**	3.6E+03	U	TSC/100 mL	P432974	
	SOP-PCR-1.3	GFD-qPCR**	3.8E+03		TSC/100 mL	P432974	
	SOP-PCR-1.4	BacR-qPCR**	3.0E+03	U	TSC/100 mL	P432974	
	SOP-PCR-1.5	DG3-qPCR**	2.6E+03	U	TSC/100 mL	P432974	
2426804	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray	1439		MPN/100 mL		

Quality Assurance Report Method Blank Results

Reference Method: SOP-PCR-1.0

Batch ID: P432974

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

Reference Method: SOP-PCR-1.2

Batch ID: P432974

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

Reference Method: SOP-PCR-1.3

Batch ID: P432974

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

Reference Method: SOP-PCR-1.4

Batch ID: P432974

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

Reference Method: SOP-PCR-1.5

Batch ID: P432974

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

Quality Assurance Report Laboratory Control Sample Accuracy

Quality Assurance Report Laboratory Control Sample Accuracy

Reference Method: SOP-PCR-1.0
Batch ID: P432974

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	80.8	99.7	P/P	50 - 175

Reference Method: SOP-PCR-1.2
Batch ID: P432974

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
GULL2-qPCR	92.9	98.8	P/P	50 - 300

Reference Method: SOP-PCR-1.3
Batch ID: P432974

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
GFD-qPCR	94.5	99.3	P/P	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P432974

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: P432974

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
DG3-qPCR	99.1	71.6	P/P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
Batch ID: P432974

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2426361	HF183-qPCR	103	109	P/P	50 - 175
2426802	HF183-qPCR	96.7		P	50 - 175
2427046	HF183-qPCR	95.4		P	50 - 175
2427265	HF183-qPCR	94.0		P	50 - 175
2427495	HF183-qPCR	92.4		P	50 - 175
2427496	HF183-qPCR	88.6		P	50 - 175
2427519	HF183-qPCR	85.8		P	50 - 175
2427528	HF183-qPCR	107		P	50 - 175
2427529	HF183-qPCR	102		P	50 - 175
2427530	HF183-qPCR	113		P	50 - 175
2427531	HF183-qPCR	113		P	50 - 175
2427532	HF183-qPCR	111		P	50 - 175
2427533	HF183-qPCR	115		P	50 - 175

Reference Method: SOP-PCR-1.2
Batch ID: P432974

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-1.2

Batch ID: P432974

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2426361	GULL2-qPCR	77.1	82.7	P/P	50 - 300
2426802	GULL2-qPCR	79.5		P	50 - 300
2427046	GULL2-qPCR	80.5		P	50 - 300
2427265	GULL2-qPCR	78.5		P	50 - 300
2427496	GULL2-qPCR	77.0		P	50 - 300
2427519	GULL2-qPCR	53.4		P	50 - 300
2427528	GULL2-qPCR	75.8		P	50 - 300
2427529	GULL2-qPCR	73.3		P	50 - 300
2427530	GULL2-qPCR	77.9		P	50 - 300
2427531	GULL2-qPCR	83.5		P	50 - 300
2427532	GULL2-qPCR	82.2		P	50 - 300
2427533	GULL2-qPCR	88.7		P	50 - 300

Reference Method: SOP-PCR-1.3

Batch ID: P432974

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2426361	GFD-qPCR	99.0	99.4	P/P	50 - 175
2426802	GFD-qPCR	95.4		P	50 - 175
2427046	GFD-qPCR	92.3		P	50 - 175
2427265	GFD-qPCR	88.5		P	50 - 175
2427496	GFD-qPCR	86.7		P	50 - 175
2427519	GFD-qPCR	90.4		P	50 - 175
2427528	GFD-qPCR	95.0		P	50 - 175
2427529	GFD-qPCR	85.0		P	50 - 175
2427530	GFD-qPCR	79.7		P	50 - 175
2427531	GFD-qPCR	77.3		P	50 - 175
2427532	GFD-qPCR	84.1		P	50 - 175
2427533	GFD-qPCR	87.8		P	50 - 175

Reference Method: SOP-PCR-1.4

Batch ID: P432974

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2426361	BacR-qPCR	102	95.4	P/P	50 - 175
2426802	BacR-qPCR	90.8		P	50 - 175
2427046	BacR-qPCR	86.3		P	50 - 175
2427265	BacR-qPCR	80.3		P	50 - 175
2427495	BacR-qPCR	69.6		P	50 - 175
2427496	BacR-qPCR	89.7		P	50 - 175
2427519	BacR-qPCR	89.8		P	50 - 175

Reference Method: SOP-PCR-1.5

Batch ID: P432974

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2426361	DG3-qPCR	105	109	P/P	50 - 175
2426802	DG3-qPCR	103		P	50 - 175
2427046	DG3-qPCR	105		P	50 - 175
2427265	DG3-qPCR	99.4		P	50 - 175
2427495	DG3-qPCR	97.7		P	50 - 175
2427496	DG3-qPCR	90.8		P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.5
Batch ID: P432974

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2427519	DG3-qPCR	92.6		P	50 - 175
2427528	DG3-qPCR	112		P	50 - 175
2427529	DG3-qPCR	110		P	50 - 175
2427530	DG3-qPCR	145		P	50 - 175
2427531	DG3-qPCR	115		P	50 - 175
2427532	DG3-qPCR	114		P	50 - 175
2427533	DG3-qPCR	120		P	50 - 175

Quality Assurance Report Precision

Reference Method: SOP-PCR-1.0
Batch ID: P432974

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2426361	HF183-qPCR	6.09	Spike	P	0 - 25

Reference Method: SOP-PCR-1.2
Batch ID: P432974

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2426361	GULL2-qPCR	6.95	Spike	P	0 - 25

Reference Method: SOP-PCR-1.3
Batch ID: P432974

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2426361	GFD-qPCR	0.487	Spike	P	0 - 25

Reference Method: SOP-PCR-1.4
Batch ID: P432974

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2426361	BacR-qPCR	6.67	Spike	P	0 - 25

Reference Method: SOP-PCR-1.5
Batch ID: P432974

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2426361	DG3-qPCR	3.90	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: 2426802
 Field Sample ID: G3SD0064

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	118	P	50 - 300
SOP-PCR-1.2	SKETA-qPCR	132	P	50 - 300
SOP-PCR-1.3	SKETA-qPCR	127	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	131	P	50 - 300
SOP-PCR-1.5	SKETA-qPCR	116	P	50 - 300

Quality Assurance Report Summary

Ref. Method	Analyte	LCS % Recovery		MS % Recovery			Precision SMP	MS
SOP-PCR-1.0	HF183-qPCR	99.7	80.8	103	109	96.7		6.09
SOP-PCR-1.2	GULL2-qPCR	98.8	92.9	77.1	82.7	79.5		6.95
SOP-PCR-1.3	GFD-qPCR	99.3	94.5	99.0	99.4	95.4		0.487
SOP-PCR-1.4	BacR-qPCR	69.2	108	102	95.4	90.8		6.67
SOP-PCR-1.5	DG3-qPCR	71.6	99.1	105	109	103		3.90

Reference Method Descriptions

Method	Description	Associated Samples
Enterolert/QT	Enterococci by Enterolert/QT by BEA South in North Port	2426803
SM 9223 Quanti-Tray	E. coli by Colilert/QT by BEA South in North Port	2426804
SOP-PCR-1.0	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	2426802
SOP-PCR-1.2	Detection and quantification of coastal bird specific Catellicoccus marimammalium Gull2 genetic marker with quantitative polymerase chain reaction	2426802
SOP-PCR-1.3	Detection and quantification of bird specific Helicobacter GFD genetic marker with quantitative polymerase chain reaction	2426802
SOP-PCR-1.4	Detection and quantification of Ruminant specific Bacteroidetes BacR genetic marker with quantitative polymerase chain reaction	2426802
SOP-PCR-1.5	Detection and quantification of canine-specific DG3 Bacteroides genetic marker with quantitative polymerase chain reaction	2426802

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
Enterolert/QT	07/25/2023	07/25/2023 13:41		07/26/2023 15:29	Benchmark EnviroAnalytical in North Port	2426803
SM 9223 Quanti-Tray	07/25/2023	07/25/2023 14:05		07/26/2023 14:57	Benchmark EnviroAnalytical in North Port	2426804
SOP-PCR-1.0	07/26/2023	07/26/2023 14:55	Claudia Castro	08/07/2023 12:37	Madeline Gruver	2426802
SOP-PCR-1.2	07/26/2023	07/26/2023 14:55	Claudia Castro	08/09/2023 14:25	Roberta Tatti	2426802
SOP-PCR-1.3	07/26/2023	07/26/2023 14:55	Claudia Castro	08/10/2023 10:08	Roberta Tatti	2426802
SOP-PCR-1.4	07/26/2023	07/26/2023 14:55	Claudia Castro	08/08/2023 15:11	Madeline Gruver	2426802
SOP-PCR-1.5	07/26/2023	07/26/2023 14:55	Claudia Castro	08/08/2023 17:19	Madeline Gruver	2426802