

Biological Analysis Report

NE-DIST-2024-02-14-02

Florida DEP Laboratory
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DOH Accreditation E31780

Overflow Analyses Performed By:
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DOH Accreditation E821059

Event Description: **Suwannee East 2024**
Request ID: **RQ-2024-02-12-11**
Customer: **NE-DIST**
Project ID: **SMP**

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Date Certified: 29-MAR-2024 14:56



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: ROCKY CREEK AT 156TH AV

Collection Date/Time: 02/13/2024 09:55

Field ID: G1NE0020

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2469047	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	2419.6		MPN/100 mL		

Sample Location: HAGUE BRANCH TRIBAT US441 80M W 77TH TERR

Collection Date/Time: 02/13/2024 10:30

Field ID: 21030152

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2469048	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	2419.6	J	MPN/100 mL		

Ref. Method and Comment:

SM 9223 B Quanti-Tray: All Quanti-Tray wells were positive. Result may be higher than reported.

Sample Location: CELLON CREEK AT SAN FELASCO SP

Collection Date/Time: 02/13/2024 11:40

Field ID: G1NE0960

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2469049	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	866.4		MPN/100 mL		

Sample Location: MILL CREEK AT OLD BELLAMY ALLIGATOR ROAD

Collection Date/Time: 02/13/2024 13:00

Field ID: 21030126

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2469050	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	2419.6	J	MPN/100 mL		
2469054	SOP-PCR-1.0	HF183-qPCR**	1.7E+03	U	TSC/100 mL	P441368	
	SOP-PCR-1.4	BacR-qPCR**	1.32E+04		TSC/100 mL	P441368	
	SOP-PCR-1.5	DG3-qPCR**	1.3E+03	U	TSC/100 mL	P441368	

Ref. Method and Comment:

SM 9223 B Quanti-Tray: All Quanti-Tray wells were positive. Result may be higher than reported.

Sample Location: PARENERS BR UPSTREAM OF CULVERTS UNDER CR1491

Collection Date/Time: 02/13/2024 13:25

Field ID: G1NE0906

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2469046	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	2419.6	J	MPN/100 mL		
2469053	SOP-PCR-1.0	HF183-qPCR**	1.7E+03	U	TSC/100 mL	P441368	
	SOP-PCR-1.4	BacR-qPCR**	2.63E+07		TSC/100 mL	P441368	
	SOP-PCR-1.5	DG3-qPCR**	1.3E+03	U	TSC/100 mL	P441368	

Ref. Method and Comment:

SM 9223 B Quanti-Tray: All Quanti-Tray wells were positive. Result may be higher than reported.

Sample Location: OLUSTEE CR AT CR 238

Collection Date/Time: 02/13/2024 14:00

Field ID: 21020009

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2469045	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	262.6		MPN/100 mL		

Quality Assurance Report
Method Blank Results

Reference Method: SOP-PCR-1.0
Batch ID: P441368

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

Reference Method: SOP-PCR-1.4
Batch ID: P441368

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

Reference Method: SOP-PCR-1.5
Batch ID: P441368

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: P441368

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	94.4	76.1	P/P	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P441368

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
BacR-qPCR	108	73.6	P/P	50 - 175

Reference Method: SOP-PCR-1.5
Batch ID: P441368

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
DG3-qPCR	94.2	52.4	P/P	50 - 175

Quality Assurance Report
Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
Batch ID: P441368

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2468556	HF183-qPCR	95.3		P	50 - 175
2468938	HF183-qPCR	94.9	96.7	P/P	50 - 175
2469053	HF183-qPCR	106		P	50 - 175
2469054	HF183-qPCR	105		P	50 - 175
2469343	HF183-qPCR	88.7		P	50 - 175
2469566	HF183-qPCR	109		P	50 - 175
2470407	HF183-qPCR	97.2		P	50 - 175
2471016	HF183-qPCR	91.2		P	50 - 175
2471167	HF183-qPCR	91.4		P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
Batch ID: P441368

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2471168	HF183-qPCR	94.8		P	50 - 175
2471252	HF183-qPCR	98.6		P	50 - 175
2471253	HF183-qPCR	90.8		P	50 - 175
2471269	HF183-qPCR	97.4		P	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P441368

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2468556	BacR-qPCR	89.4		P	50 - 175
2468938	BacR-qPCR	105	96.1	P/P	50 - 175
2469053	BacR-qPCR	84.7		P	50 - 175
2469054	BacR-qPCR	81.3		P	50 - 175
2469343	BacR-qPCR	72.9		P	50 - 175
2469566	BacR-qPCR	75.6		P	50 - 175
2470407	BacR-qPCR	91.5		P	50 - 175
2471016	BacR-qPCR	77.1		P	50 - 175
2471167	BacR-qPCR	79.4		P	50 - 175
2471168	BacR-qPCR	82.7		P	50 - 175
2471252	BacR-qPCR	84.6		P	50 - 175
2471253	BacR-qPCR	68.9		P	50 - 175
2471269	BacR-qPCR	82.0		P	50 - 175

Reference Method: SOP-PCR-1.5
Batch ID: P441368

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2468556	DG3-qPCR	91.5		P	50 - 175
2468938	DG3-qPCR	94.1	97.2	P/P	50 - 175
2469053	DG3-qPCR	94.7		P	50 - 175
2469054	DG3-qPCR	94.2		P	50 - 175
2469343	DG3-qPCR	88.1		P	50 - 175
2469566	DG3-qPCR	92.8		P	50 - 175
2470407	DG3-qPCR	90.6		P	50 - 175
2471016	DG3-qPCR	91.1		P	50 - 175
2471167	DG3-qPCR	95.1		P	50 - 175
2471168	DG3-qPCR	94.9		P	50 - 175
2471252	DG3-qPCR	89.6		P	50 - 175
2471253	DG3-qPCR	79.3		P	50 - 175
2471269	DG3-qPCR	92.0		P	50 - 175

Quality Assurance Report Precision

Reference Method: SOP-PCR-1.0
Batch ID: P441368

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2468938	HF183-qPCR	1.80	Spike	P	0 - 25

Quality Assurance Report

Precision

Reference Method: SOP-PCR-1.4

Batch ID: P441368

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2468938	BacR-qPCR	9.20	Spike	P	0 - 25

Reference Method: SOP-PCR-1.5

Batch ID: P441368

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2468938	DG3-qPCR	3.18	Spike	P	0 - 25

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

Quality Assurance Report

Surrogates

Lab Sample ID: 2469053

Field Sample ID: G1NE0906

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	101	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	118	P	50 - 300
SOP-PCR-1.5	SKETA-qPCR	96.9	P	50 - 300

Lab Sample ID: 2469054

Field Sample ID: 21030126

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	102	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	122	P	50 - 300
SOP-PCR-1.5	SKETA-qPCR	95.7	P	50 - 300

Quality Assurance Report

Summary

Ref. Method	Analyte	LCS % Recovery		MS % Recovery			Precision	
							LCS	MS
SOP-PCR-1.0	HF183-qPCR	76.1	94.4	95.3	94.9	96.7	Precision SMP	1.80
SOP-PCR-1.4	BacR-qPCR	108	73.6	89.4	105	96.1		9.20
SOP-PCR-1.5	DG3-qPCR	52.4	94.2	91.5	94.1	97.2		3.18

Reference Method Descriptions

Method	Description	Associated Samples
SM 9223 B Quanti-Tray	E. coli by Colilert/QT by Diversified Environmental Laboratories in Jacksonville	2469045, 2469046, 2469047, 2469048, 2469049, 2469050
SOP-PCR-1.0	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	2469053, 2469054

Reference Method Descriptions

Method	Description	<u>Associated Samples</u>
SOP-PCR-1.4	Detection and quantification of Ruminant specific Bacteroidetes BacR genetic marker with quantitative polymerase chain reaction	2469053, 2469054
SOP-PCR-1.5	Detection and quantification of canine-specific DG3 Bacteroides genetic marker with quantitative polymerase chain reaction	2469053, 2469054

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	02/13/2024	02/13/2024 16:07		02/14/2024 10:32	Diversified-Jacksonville	2469045, 2469046, 2469047, 2469048, 2469049, 2469050
SOP-PCR-1.0	02/14/2024	02/14/2024 13:16	Lily Gregson	03/01/2024 08:48	Roberta Tatti	2469053, 2469054
SOP-PCR-1.4	02/14/2024	02/14/2024 13:16	Lily Gregson	03/04/2024 08:51	Roberta Tatti	2469053, 2469054
SOP-PCR-1.5	02/14/2024	02/14/2024 13:16	Lily Gregson	03/13/2024 08:53	Roberta Tatti	2469053, 2469054