

Biological Analysis Report

NW-DIST-2022-07-21-01

Florida Department of Environmental Protection
Central Laboratory
2600 Blair Stone Road
Tallahassee, FL 32399-2400
DOH Accreditation E31780

Event Description: **SMP/TREND RUN**
Request ID: **RQ-2022-07-18-39**
Customer: **NW-DIST**
Project ID: **SMP**

Send Reports to:
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160 W Government Street
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Certified by: Sarah Menz, Environmental Administrator

Date Certified: 29-AUG-2022 10:54

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 group are included in this report: 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: Yellow River at Oak Grove Boat Ramp

Collection Date/Time: 07/20/2022 11:05

Field ID: G4NW0526

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2343224	SOP-PCR-1.0	HF183-qPCR**	3.3E+03	U	TSC/100 mL	P416899	
	SOP-PCR-1.4	BacR-qPCR**	4.57E+05		TSC/100 mL	P416899	MS
	SOP-PCR-1.5	DG3-qPCR**	2.6E+03	U	TSC/100 mL	P416899	

Quality Assurance Report Method Blank Results

Reference Method: SOP-PCR-1.0

Batch ID: P416899

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

Reference Method: SOP-PCR-1.4

Batch ID: P416899

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

Reference Method: SOP-PCR-1.5

Batch ID: P416899

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: P416899

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	95.4	79.4	P/P	50 - 175

Reference Method: SOP-PCR-1.4

Batch ID: P416899

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
BacR-qPCR	98.8	75.5	P/P	50 - 175

Reference Method: SOP-PCR-1.5

Batch ID: P416899

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
DG3-qPCR	75.1	78.6	P/P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0

Batch ID: P416899

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2342444	HF183-qPCR	50.6	59.3	P/P	50 - 175
2342735	HF183-qPCR	99.8		P	50 - 175
2342736	HF183-qPCR	88.7		P	50 - 175
2343219	HF183-qPCR	90.5		P	50 - 175
2343224	HF183-qPCR	86.5		P	50 - 175
2344365	HF183-qPCR	73.1		P	50 - 175
2344366	HF183-qPCR	72.5		P	50 - 175
2344708	HF183-qPCR	94.2		P	50 - 175
2344763	HF183-qPCR	128		P	50 - 175
2344764	HF183-qPCR	83.9		P	50 - 175
2345085	HF183-qPCR	89.4		P	50 - 175
2345123	HF183-qPCR	92.2		P	50 - 175

Reference Method: SOP-PCR-1.4

Batch ID: P416899

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2342444	BacR-qPCR	91.1	91.2	P/P	50 - 175
2342735	BacR-qPCR	69.7		P	50 - 175
2342736	BacR-qPCR	83.1		P	50 - 175
2343219	BacR-qPCR	74.8		P	50 - 175
2343224	BacR-qPCR	59.0		P	50 - 175
2344365	BacR-qPCR	47.3		F	50 - 175
2344366	BacR-qPCR	52.3		P	50 - 175
2344708	BacR-qPCR	81.9		P	50 - 175
2344763	BacR-qPCR	81.3		P	50 - 175
2344764	BacR-qPCR	84.8		P	50 - 175
2345085	BacR-qPCR	78.2		P	50 - 175
2345123	BacR-qPCR	82.2		P	50 - 175

Reference Method: SOP-PCR-1.5

Batch ID: P416899

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2342444	DG3-qPCR	97.6	97.9	P/P	50 - 175
2342735	DG3-qPCR	71.7		P	50 - 175
2342736	DG3-qPCR	93.9		P	50 - 175
2343219	DG3-qPCR	99.7		P	50 - 175
2343224	DG3-qPCR	94.8		P	50 - 175
2344365	DG3-qPCR	67.1		P	50 - 175
2344366	DG3-qPCR	71.7		P	50 - 175
2344708	DG3-qPCR	84.9		P	50 - 175
2344763	DG3-qPCR	86.3		P	50 - 175
2344764	DG3-qPCR	89.2		P	50 - 175
2345085	DG3-qPCR	88.4		P	50 - 175
2345123	DG3-qPCR	93.9		P	50 - 175

Quality Assurance Report Precision

Quality Assurance Report Precision

Reference Method: SOP-PCR-1.0
 Batch ID: P416899

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2342444	HF183-qPCR	5.93	Spike	P	0 - 25

Reference Method: SOP-PCR-1.4
 Batch ID: P416899

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2342444	BacR-qPCR	0.178	Spike	P	0 - 25

Reference Method: SOP-PCR-1.5
 Batch ID: P416899

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2342444	DG3-qPCR	0.363	Spike	P	0 - 25

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

Quality Assurance Report Surrogates

Lab Sample ID: 2343224
 Field Sample ID: G4NW0526

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	78.0	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	113	P	50 - 300
SOP-PCR-1.5	SKETA-qPCR	79.0	P	50 - 300

Quality Assurance Report Summary

Ref. Method	Analyte	LCS % Recovery		MS % Recovery			Precision SMP	MS
SOP-PCR-1.0	HF183-qPCR	79.4	95.4	50.6	59.3	99.8		5.93
				88.7				
SOP-PCR-1.4	BacR-qPCR	75.5	98.8	91.1	91.2	83.1		0.178
				69.7				
SOP-PCR-1.5	DG3-qPCR	78.6	75.1	97.6	97.9	71.7		0.363
				93.9				

Reference Method Descriptions

Method	Description	Associated Samples
SOP-PCR-1.0	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	2343224
SOP-PCR-1.4	Detection and quantification of Ruminant specific Bacteroidetes BacR genetic marker with quantitative polymerase chain reaction	2343224

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	2343224

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
SOP-PCR-1.0	07/21/2022	07/21/2022 14:10	Claudia Castro	08/10/2022 14:26	DeAsia Armster	2343224
SOP-PCR-1.4	07/21/2022	07/21/2022 14:10	Claudia Castro	08/11/2022 10:27	Roberta Tatti	2343224
SOP-PCR-1.5	07/21/2022	07/21/2022 14:10	Claudia Castro	08/11/2022 12:48	DeAsia Armster	2343224