

Biological Analysis Report

CEN-DIST-2019-10-01-01

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

Overflow Analyses Performed By:
Advanced Environmental Laboratories, Inc.
380 North Lake Blvd, Ste 1048
Altamonte Springs, FL 32701
DOH Accreditation E53076

Event Description: **UnBranch X2 / Groover X2**
Request ID: **RQ-2019-09-02-67**
Customer: **CEN-DIST**
Project ID: **SMP**

Send Reports to:
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Certified by: Cheryl A. Swanson, Program Administrator

Date Certified: 18-OCT-2019 15:58



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: Groover Branch at Tymber Run Road

Collection Date/Time: 09/30/2019 11:58

Field ID: G5CE0041

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2123783	EPA 1603	Escherichia coli Membrane Filter	330		cfu/100 mL		

Sample Location: Groover Branch at Airport Road Bridge

Collection Date/Time: 09/30/2019 12:11

Field ID: G5CE0040

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2123784	EPA 1603	Escherichia coli Membrane Filter	120	B	cfu/100 mL		

Sample Location: Unnamed Branch @ 60m upstream of Airport Road

Collection Date/Time: 09/30/2019 11:13

Field ID: G5CE0084

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2123781	EPA 1603	Escherichia coli Membrane Filter	570		cfu/100 mL		
2123785	SOP-PCR-1.0	HF183-qPCR**	140	U	GEU/100 mL	P371626	

Ref. Method and Comment:
 SOP-PCR-1.0: Precision data unavailable for Analysis Batch A95025.

Sample Location: Unnamed Branch @ Bear Creek Path

Collection Date/Time: 09/30/2019 11:27

Field ID: G5CE0032

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2123782	EPA 1603	Escherichia coli Membrane Filter	980	B	cfu/100 mL		
2123786	SOP-PCR-1.0	HF183-qPCR**	140	U	GEU/100 mL	P371626	

Quality Assurance Report Method Blank Results

Reference Method: SOP-PCR-1.0
 Batch ID: P371626

Component	Result	Code	Units
HF183-qPCR	280	U	GEU/100 mL
HF183-qPCR	280	U	GEU/100 mL

Quality Assurance Report Laboratory Control Sample Accuracy

Reference Method: SOP-PCR-1.0
 Batch ID: P371626

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	151	90.1	P/P	50 - 175
HF183-qPCR	148	100	P/P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
 Batch ID: P371626

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2123785	HF183-qPCR	101		P	50 - 175
2123786	HF183-qPCR	117	112	P/P	50 - 175
2123983	HF183-qPCR	119		P	50 - 175
2123984	HF183-qPCR	114		P	50 - 175
2123985	HF183-qPCR	121		P	50 - 175
2124016	HF183-qPCR	100		P	50 - 175
2124017	HF183-qPCR	97.1		P	50 - 175
2124376	HF183-qPCR	114		P	50 - 175
2124432	HF183-qPCR	114		P	50 - 175
2124433	HF183-qPCR	110		P	50 - 175
2124434	HF183-qPCR	114		P	50 - 175
2124578	HF183-qPCR	90.9		P	50 - 175
2124677	HF183-qPCR	94.5		P	50 - 175
2124678	HF183-qPCR	92.0		P	50 - 175

Quality Assurance Report Precision

Reference Method: SOP-PCR-1.0
 Batch ID: P371626

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2123786	HF183-qPCR	5.02	Spike	P	0 - 25

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

Quality Assurance Report Surrogates

Lab Sample ID: 2123785
 Field Sample ID: G5CE0084

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	84.2	P	50 - 300

Lab Sample ID: 2123786
 Field Sample ID: G5CE0032

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	91.2	P	50 - 300

Quality Assurance Report Summary

Ref. Method	Analyte	LCS % Recovery			MS % Recovery			LCS	Precision SMP	MS
SOP-PCR-1.0	HF183-qPCR	151	100	148	117	112	119			5.02
		90.1			114					

Reference Method Descriptions

Method / DoH Cert #	Description	<u>Associated Samples</u>
EPA 1603 / E53076	E. coli by MF by AEL in Altamonte Springs	2123781, 2123782, 2123783, 2123784
SOP-PCR-1.0 / E31780	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	2123785, 2123786

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
EPA 1603	09/30/2019	09/30/2019 15:55		10/01/2019 15:49	Advanced Environmental Laboratory	2123781, 2123782, 2123783, 2123784
SOP-PCR-1.0	10/01/2019	10/01/2019 15:45	Angela Smith	10/14/2019 12:26	Lauren Campbell	2123786
	10/01/2019	10/01/2019 15:45	Angela Smith	10/16/2019 12:17	Lauren Campbell	2123785