

Biological Analysis Report

SW-DIST-2019-10-22-01

Florida Department of Environmental Protection
Central Laboratory
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DOH Accreditation E31780

Overflow Analyses Performed By:
Advanced Environmental Laboratories - Tampa
9610 Princess Palm Avenue
Tampa, Florida 33619
DOH Accreditation E84589

Event Description: **English, Pemberton, Baker, Spartman, Mustang, Poley**
Request ID: **RQ-2019-10-21-05**
Customer: **SW-DIST**
Project ID: **SMP**

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Date Certified: 15-NOV-2019 15:28



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: Poley Creek @ SR 60

Collection Date/Time: 10/21/2019 11:10

Field ID: G2SW0049

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2130284	SOP-PCR-1.0	HF183-qPCR**	140	U	GEU/100 mL	P372898	
	SOP-PCR-1.4	BacR-qPCR**	5.9E+03	I	TSC/100 mL	P372898	LCS
2130290	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	521		MPN/100 mL		

Ref. Method and Comment:
 SOP-PCR-1.0: Precision data unavailable due to inhibition in QC sample.

Sample Location: Spartman Branch at Harvey Tew Rd

Collection Date/Time: 10/21/2019 10:20

Field ID: G2SW0067

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2130285	SOP-PCR-1.0	HF183-qPCR**	140	U	GEU/100 mL	P372898	
	SOP-PCR-1.4	BacR-qPCR**	3.38E+04		TSC/100 mL	P372898	LCS
2130291	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	333		MPN/100 mL		

Ref. Method and Comment:
 SOP-PCR-1.0: Precision data unavailable due to inhibition in QC sample.

Sample Location: Pemberton Creek @ Gallagher Rd

Collection Date/Time: 10/21/2019 10:05

Field ID: G2SW0085

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2130293	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	582		MPN/100 mL		

Sample Location: Mustang Ranch Creek @ Hopewell

Collection Date/Time: 10/21/2019 10:50

Field ID: G2SW0026

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2130286	SOP-PCR-1.0	HF183-qPCR**	140	U	GEU/100 mL	P372898	
	SOP-PCR-1.4	BacR-qPCR**	1.5E+03	U	TSC/100 mL	P372898	LCS
2130292	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	297		MPN/100 mL		

Ref. Method and Comment:
 SOP-PCR-1.0: Precision data unavailable due to inhibition in QC sample.

Quality Assurance Report Method Blank Results

Reference Method: SOP-PCR-1.0
Batch ID: P372898

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

Reference Method: SOP-PCR-1.4
Batch ID: P372898

Component	Result	Code	Units
BacR-qPCR	3.0E+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: P372898

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	79.3	152	P/P	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P372898

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
BacR-qPCR	117	304	P/F	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
Batch ID: P372898

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2130156	HF183-qPCR	110		P	50 - 175
2130207	HF183-qPCR	105		P	50 - 175
2130284	HF183-qPCR	97.8		P	50 - 175
2130285	HF183-qPCR	108		P	50 - 175
2130286	HF183-qPCR	97.8		P	50 - 175
2130342	HF183-qPCR	108		P	50 - 175
2130343	HF183-qPCR	103		P	50 - 175
2130344	HF183-qPCR	109		P	50 - 175
2130581	HF183-qPCR	95.0		P	50 - 175
2130582	HF183-qPCR	97.0		P	50 - 175
2130583	HF183-qPCR	101		P	50 - 175
2130632	HF183-qPCR	112		P	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P372898

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2130156	BacR-qPCR	90.3		P	50 - 175
2130207	BacR-qPCR	103		P	50 - 175
2130284	BacR-qPCR	70.1		P	50 - 175
2130285	BacR-qPCR	98.2		P	50 - 175
2130286	BacR-qPCR	78.1		P	50 - 175
2130581	BacR-qPCR	72.4		P	50 - 175
2130582	BacR-qPCR	80.1		P	50 - 175
2130583	BacR-qPCR	79.0		P	50 - 175
2132982	BacR-qPCR	68.7	60.9	P/P	50 - 175

Quality Assurance Report Precision

Reference Method: SOP-PCR-1.4
Batch ID: P372898

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2132982	BacR-qPCR	12.0	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: 2130284
Field Sample ID: G2SW0049

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	93.2	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	101	P	50 - 300

Lab Sample ID: 2130285
Field Sample ID: G2SW0067

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	86.8	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	89.5	P	50 - 300

Lab Sample ID: 2130286
Field Sample ID: G2SW0026

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	84.7	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	93.7	P	50 - 300

Quality Assurance Report Summary

Ref. Method	Analyte	LCS % Recovery		MS % Recovery			LCS	Precision SMP	MS
SOP-PCR-1.0	HF183-qPCR	152	79.3	110	105	97.8			
				108					
SOP-PCR-1.4	BacR-qPCR	304	117	90.3	103	70.1			12.0
				98.2					

Reference Method Descriptions

Method / DoH Cert #	Description	Associated Samples
SM 9223 B Quanti-Tray / E84589	E. coli by Colilert-18/QT by AEL in Tampa	2130290, 2130291, 2130292, 2130293
SOP-PCR-1.0 / E31780	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	2130284, 2130285, 2130286
SOP-PCR-1.4 / E31780	Detection and quantification of Ruminant specific Bacteroidetes BacR genetic marker with quantitative polymerase chain reaction	2130284, 2130285, 2130286

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	10/21/2019	10/21/2019 12:38		10/22/2019 08:35	AEL - Tampa	2130290, 2130291, 2130292, 2130293
SOP-PCR-1.0	10/22/2019	10/22/2019 13:40	Angela Smith	11/12/2019 11:27	Lauren Campbell	2130284, 2130285, 2130286
SOP-PCR-1.4	10/22/2019	10/22/2019 13:40	Angela Smith	11/13/2019 14:38	Lauren Campbell	2130284, 2130285, 2130286