

Biological Analysis Report

SW-DIST-2019-05-07-01

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

Event Description: **Bishop Creek-Pinellas Sites 12-04, 12-02**

Request ID: **RQ-2019-05-06-06**

Customer: **SW-DIST**

Project ID: **SMP**

Send Reports to:

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Date Certified: 17-JUN-2019 14:28



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 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: BISHOP CRK @ HARBOR LAKES

Collection Date/Time: 05/06/2019 10:02

Field ID: G1SW0061

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2083596	SOP-PCR-4.0	HF183 - Human Source Marker**	Below Threshold			P363169	

Ref. Method and Comment:
SOP-PCR-4.0: Bacterial results for threshold obtained from county lab.

Sample Location: BISHOP CRK @ OAKCREST DR.

Collection Date/Time: 05/06/2019 10:38

Field ID: G1SW0063

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2083597	SOP-PCR-1.0	HF183-qPCR**	94	U	GEU/100 mL	P363934	LCS
	SOP-PCR-1.2	GULL2-qPCR**	1.1E+04	U	TSC/100 mL	P363934	LCS
	SOP-PCR-1.3	GFD-purified-qPCR**	110	UJ	TSC/100 mL	P365054	MS
	SOP-PCR-1.4	BacR-qPCR**	1.0E+03	U	TSC/100 mL	P363934	
	SOP-PCR-4.0	HF183 - Human Source Marker**	Above Threshold			P363169	

Ref. Method and Comment:
SOP-PCR-1.2: Precision data unavailable for Analysis Batch A92002
SOP-PCR-1.3: Quality control failure(s) observed. See QA report for details.
SOP-PCR-1.4: Precision data unavailable for Analysis Batch A91985
SOP-PCR-4.0: Bacterial results for threshold obtained from county lab.

Quality Assurance Report Method Blank Results

Reference Method: SOP-PCR-1.0
Batch ID: P363934

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

Reference Method: SOP-PCR-1.2
Batch ID: P363934

Component	Result	Code	Units
GULL2-qPCR	1.7E+04	U	TSC/100 mL

Reference Method: SOP-PCR-1.3
Batch ID: P365054

Component	Result	Code	Units
GFD-purified-qPCR	340	U	TSC/100 mL

Reference Method: SOP-PCR-1.4
Batch ID: P363934

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: P363934

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	186	115	F/P	50 - 175
HF183-qPCR	164	153	P/P	50 - 175

Reference Method: SOP-PCR-1.2
Batch ID: P363934

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
GULL2-qPCR	184	435	P/F	50 - 300

Reference Method: SOP-PCR-1.3
Batch ID: P365054

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
GFD-purified-qPCR	96.7	85.2	P/P	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P363934

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
BacR-qPCR	81.7	76.1	P/P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
Batch ID: P363934

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2083344	HF183-qPCR	128		P	50 - 175
2083597	HF183-qPCR	105		P	50 - 175
2085191	HF183-qPCR	106		P	50 - 175
2085852	HF183-qPCR	129		P	50 - 175
2085853	HF183-qPCR	111		P	50 - 175
2085951	HF183-qPCR	106		P	50 - 175
2085952	HF183-qPCR	155		P	50 - 175
2085953	HF183-qPCR	106		P	50 - 175
2085954	HF183-qPCR	133	119	P/P	50 - 175
2086433	HF183-qPCR	122		P	50 - 175
2086434	HF183-qPCR	120		P	50 - 175
2086435	HF183-qPCR	110		P	50 - 175
2090303	HF183-qPCR	66.4		P	50 - 175

Reference Method: SOP-PCR-1.2
Batch ID: P363934

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2083344	GULL2-qPCR	119		P	50 - 300
2083597	GULL2-qPCR	111		P	50 - 300
2085191	GULL2-qPCR	105		P	50 - 300
2085852	GULL2-qPCR	103		P	50 - 300
2085853	GULL2-qPCR	100		P	50 - 300

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.3

Batch ID: P365054

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2083344	GFD-purified-qPCR	187		F	50 - 175
2083597	GFD-purified-qPCR	198		F	50 - 175
2085191	GFD-purified-qPCR	162		P	50 - 175
2085852	GFD-purified-qPCR	198		F	50 - 175
2085853	GFD-purified-qPCR	160		P	50 - 175
2091364	GFD-purified-qPCR	235		F	50 - 175
2091365	GFD-purified-qPCR	220	208	F/F	50 - 175
2091366	GFD-purified-qPCR	219		F	50 - 175
2091463	GFD-purified-qPCR	171		P	50 - 175

Reference Method: SOP-PCR-1.4

Batch ID: P363934

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2083344	BacR-qPCR	68.1		P	50 - 175
2083597	BacR-qPCR	50.8		P	50 - 175
2085191	BacR-qPCR	55.7		P	50 - 175
2085852	BacR-qPCR	72.9		P	50 - 175
2085853	BacR-qPCR	56.6		P	50 - 175
2090303	BacR-qPCR	58.2		P	50 - 175

Quality Assurance Report Precision

Reference Method: SOP-PCR-1.0

Batch ID: P363934

Replicated

Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2085954	HF183-qPCR	11.2	Spike	P	0 - 25
2091365	HF183-qPCR	9.32	Spike	P	0 - 25

Reference Method: SOP-PCR-1.3

Batch ID: P365054

Replicated

Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2091365	GFD-purified-qPCR	6.49	Spike	P	0 - 25

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

Replicate spike precision may be reported when sample results are below quantifiable levels.

Quality Assurance Report Surrogates

Lab Sample ID: 2083597

Field Sample ID: G1SW0063

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	80.4	P	50 - 300
SOP-PCR-1.2	SKETA-qPCR	123	P	50 - 300

Quality Assurance Report Surrogates

Lab Sample ID: 2083597
Field Sample ID: G1SW0063

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.4	SKETA-qPCR	96.0	P	50 - 300

Quality Assurance Report Summary

Ref. Method	Analyte	LCS % Recovery			MS % Recovery			LCS	Precision SMP	MS
SOP-PCR-1.0	HF183-qPCR	186	153	164	106	155	106		11.2	9.32
		115			133					
SOP-PCR-1.2	GULL2-qPCR	435	184		119	105	103			
					100					
SOP-PCR-1.3	GFD-purified-qPCR	85.2	96.7		235	220	208		6.49	
					219					
SOP-PCR-1.4	BacR-qPCR	76.1	81.7		68.1	55.7	72.9			
					58.2					

Reference Method Descriptions

Method / DoH Cert #	Description	Associated Samples
SOP-PCR-1.0 / E31780	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	2083597
SOP-PCR-1.2 / E31780	Detection and quantification of coastal bird specific Catellicoccus marimammalium Gull2 genetic marker with quantitative polymerase chain reaction	2083597
SOP-PCR-1.3 / E31780	Detection and quantification of bird specific Helicobacter GFD genetic marker with quantitative polymerase chain reaction	2083597
SOP-PCR-1.4 / E31780	Detection and quantification of Ruminant specific Bacteroidetes BacR genetic marker with quantitative polymerase chain reaction	2083597
SOP-PCR-4.0 / E31780	Preparation of Samples by Filtration and held for qPCR Analysis	2083596, 2083597

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
SOP-PCR-1.0	05/07/2019	05/07/2019 15:15	Angela Smith	06/06/2019 12:17	Mailin Lopez	2083597
SOP-PCR-1.2	05/07/2019	05/07/2019 15:15	Angela Smith	06/11/2019 08:29	Mailin Lopez	2083597
SOP-PCR-1.3	05/07/2019	05/07/2019 15:15	Angela Smith	06/12/2019 11:50	Mailin Lopez	2083597
SOP-PCR-1.4	05/07/2019	05/07/2019 15:15	Angela Smith	06/10/2019 13:07	Mailin Lopez	2083597
SOP-PCR-4.0	05/07/2019	05/07/2019 15:15	Angela Smith	06/05/2019 10:34	Sarah Menz	2083596, 2083597