

Biological Analysis Report

CEN-DIST-2017-06-16-03

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

Overflow Analyses Performed By:
Flowers Chemical Laboratories Inc. - Altamonte
Springs
481 Newburyport Avenue
Altamonte Springs, Florida 32701
DOH Accreditation E83018

Event Description: **Rose / Strickland / UnBranch 2642**
Request ID: **RQ-2017-06-05-38**
Customer: **CEN-DIST**
Project ID: **SMP**

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Certified by: Cheryl A. Swanson, Environmental Manager

Date Certified: 13-SEP-2017 07:59



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: Chlorophyll/Grain Size/BOD, 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 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: STRICKLAND BAY @ 100M SW OF US1 BRIDGE

Collection Date/Time: 06/15/2017 12:43

Field ID: G5CE0090

Matrix: W-SURF-SLT

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1907077	SM 10200 H (mod.)	Chlorophyll-a, Corrected	15		ug/L	P327246	
		Phaeophytin-a	3.7		ug/L	P327246	
		Chlorophyll/Phaeophytin Ratio	1.6			P327246	
		Chlorophyll-a, Uncorrected	18		ug/L	P327246	

Sample Location: STRICKLAND BAY @ 100M SW OF US1 BRIDGE

Collection Date/Time: 06/15/2017 12:43

Field ID: G5CE0090

Matrix: W-SURF-SLT

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1907076	Enterolert/QT	Enterococci-Quanti-Tray	72.8		MPN/100 mL		
1907080	SOP-PCR-1.0	HF183-qPCR**	71	U	GEU/100 mL	P326834	
	SOP-PCR-1.2	GULL2-qPCR**	1.3E+03	T	TSC/100 mL	P326971	RPD
	SOP-PCR-1.3	GFD-purified-qPCR**	16	TJ	TSC/100 mL	P326776	LCS, MS, RPD
	SOP-PCR-4.0	HF183 - Human Source Marker**	Below Threshold				

Ref. Method and Comment:

SOP-PCR-1.0: No amplification of HF183 marker.

SOP-PCR-1.3: Quality control failure(s) observed. See QA report for details.

Quality Assurance Report Method Blank Results

Reference Method: SM 10200 H (mod.)

Batch ID: P327246

Component	Result	Code	Units
Chlorophyll-a, Corrected	0.55	U	ug/L
Chlorophyll-a, Uncorrected	0.40	U	ug/L
Phaeophytin-a	0.40	U	ug/L

Reference Method: SOP-PCR-1.0

Batch ID: P326834

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

Reference Method: SOP-PCR-1.2

Batch ID: P326971

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

Reference Method: SOP-PCR-1.3

Batch ID: P326776

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

Quality Assurance Report Laboratory Control Sample Accuracy

Quality Assurance Report Laboratory Control Sample Accuracy

Reference Method: SM 10200 H (mod.)
 Batch ID: P327246

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	97.7		P	84 - 116

Reference Method: SOP-PCR-1.0
 Batch ID: P326834

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	79.4	67.8	P/P	50 - 175

Reference Method: SOP-PCR-1.2
 Batch ID: P326971

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
GULL2-qPCR	81.6	112	P/P	50 - 300

Reference Method: SOP-PCR-1.3
 Batch ID: P326776

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
GFD-purified-qPCR	56.6	23.7	P/F	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
 Batch ID: P326834

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
1886719	HF183-qPCR	114		P	50 - 175
1893375	HF183-qPCR	83.0		P	50 - 175
1895760	HF183-qPCR	82.6		P	50 - 175
1896808	HF183-qPCR	91.1		P	50 - 175
1906648	HF183-qPCR	80.5	83.3	P/P	50 - 175
1906649	HF183-qPCR	84.4		P	50 - 175
1906650	HF183-qPCR	80.0		P	50 - 175
1906651	HF183-qPCR	77.5		P	50 - 175
1907080	HF183-qPCR	85.0		P	50 - 175

Reference Method: SOP-PCR-1.2
 Batch ID: P326971

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
1907080	GULL2-qPCR	50.8		P	50 - 300

Reference Method: SOP-PCR-1.3
 Batch ID: P326776

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
1905391	GFD-purified-qPCR	96.0		P	50 - 175
1905392	GFD-purified-qPCR	109		P	50 - 175
1905393	GFD-purified-qPCR	138		P	50 - 175
1905394	GFD-purified-qPCR	93.0		P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.3
Batch ID: P326776

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
1905395	GFD-purified-qPCR	28.7		F	50 - 175
1905396	GFD-purified-qPCR	71.1		P	50 - 175
1905993	GFD-purified-qPCR	73.9	173	P/P	50 - 175
1906939	GFD-purified-qPCR	18.7		F	50 - 175
1907080	GFD-purified-qPCR	20.5		F	50 - 175
1908580	GFD-purified-qPCR	77.1		P	50 - 175

Quality Assurance Report Precision

Reference Method: SM 10200 H (mod.)
Batch ID: P327246

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
1906992	Chlorophyll/Phaeophytin Ratio	0.0	Sample	P	0 - 25
1906992	Chlorophyll-a, Corrected	0.646	Sample	P	0 - 25
1906992	Chlorophyll-a, Uncorrected	3.26	Sample	P	0 - 25
1906992	Phaeophytin-a	16.0	Sample	P	0 - 25

Reference Method: SOP-PCR-1.0
Batch ID: P326834

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
1906648	HF183-qPCR	3.52	Spike	P	0 - 25

Reference Method: SOP-PCR-1.2
Batch ID: P326971

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
1907519	GULL2-qPCR	124	Spike	F	0 - 25

Reference Method: SOP-PCR-1.3
Batch ID: P326776

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
1905993	GFD-purified-qPCR	80.5	Spike	F	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: 1907080
Field Sample ID: G5CE0090

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
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Quality Assurance Report Surrogates

Lab Sample ID: 1907080
Field Sample ID: G5CE0090

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	107	P	50 - 300
SOP-PCR-1.2	SKETA-qPCR	121	P	50 - 300

Quality Assurance Report Summary

Ref. Method	Analyte	LCS % Recovery		MS % Recovery			LCS	Precision SMP	MS
SM 10200 H (mod.)	Chlorophyll/Phaeophytin Ratio							0.0	
	Chlorophyll-a, Corrected	97.7						0.646	
	Chlorophyll-a, Uncorrected							3.26	
	Phaeophytin-a							16.0	
SOP-PCR-1.0	HF183-qPCR	67.8	79.4	114	83.0	82.6			3.52
				91.1					
SOP-PCR-1.2	GULL2-qPCR	112	81.6	50.8					124
SOP-PCR-1.3	GFD-purified-qPCR	23.7	56.6	73.9	173	18.7			80.5
				20.5					

Reference Method Descriptions

Method / DoH Cert #	Description	Associated Samples
Enterolert/QT / E83018	Enterococci by Enterolert Quanti-Tray method by Flowers Overflow Laboratory for Central District	1907076
SM 10200 H (mod.) / E31780	Phytoplankton chlorophyll-a corrected, uncorrected, phaeophytin and ratio by spectrophotometry	1907077
SOP-PCR-1.0 / E31780	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	1907080
SOP-PCR-1.2 / E31780	Detection and quantification of coastal bird specific Catellicoccus marimammalium genetic marker with quantitative polymerase chain reaction	1907080
SOP-PCR-1.3 / E31780	Detection and quantification of bird specific Helicobacter genetic marker with quantitative polymerase chain reaction	1907080
SOP-PCR-4.0 / E31780	Preparation of Samples by Filtration and held for qPCR Analysis	1907080

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
Enterolert/QT	06/15/2017			06/15/2017 16:45	Flowers - Altamonte Springs	1907076
SM 10200 H (mod.)	06/16/2017	06/16/2017 13:00	Morgan Lane	06/21/2017	Kim M. Spencer	1907077
SOP-PCR-1.0	06/16/2017	06/16/2017 13:45	Puja Jasrotia	06/19/2017	Puja Jasrotia	1907080
SOP-PCR-1.2	06/16/2017	06/16/2017 13:45	Puja Jasrotia	07/05/2017	Sarah Menz	1907080
SOP-PCR-1.3	06/16/2017	06/16/2017 13:45	Sarah Menz	06/27/2017	Sarah Menz	1907080
SOP-PCR-4.0	06/16/2017	06/16/2017 13:45	Avril Wood	07/14/2017	Puja Jasrotia	1907080