

Biological Analysis Report

SO-DIST-2018-04-05-01

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

Overflow Analyses Performed By:
Sanders Laboratories
10090 Bavaria Road
Ft. Myers, FL 33913
DOH Accreditation E85457

Event Description: **Sarasota Streams**
Request ID: **RQ-2018-04-02-67**
Customer: **SO-DIST**
Project ID: **SMP**

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Date Certified: 29-MAY-2018 12:34



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: Phillippi Creek@Proctor

Collection Date/Time: 04/04/2018 10:30

Field ID: G3SD0091

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1981567	SM 10200 H (mod.)	Chlorophyll-a, Corrected	1.2	I	ug/L	P343380	
		Phaeophytin-a	1.4	I	ug/L	P343380	
		Chlorophyll-a, Uncorrected	2.0	I	ug/L	P343380	

Ref. Method and Comment:

SM 10200 H (mod.): Precision data is not available for at least one component due to the small amount of analyte in the QC sample. Refer to QA report for available precision data.

Sample Location: Phillippi Creek@Proctor

Collection Date/Time: 04/04/2018 10:30

Field ID: G3SD0091

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1981573	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray**	325.5		MPN/100 mL		

Sample Location: Matheny Creek @BenevaRd

Collection Date/Time: 04/04/2018 11:00

Field ID: G3SD0092

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1981574	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray**	46.4		MPN/100 mL		
1981664	SOP-PCR-4.0	HF183 - Human Source Marker**	Below Threshold				

Sample Location: North Creek(Tidal)@US41

Collection Date/Time: 04/04/2018 11:30

Field ID: G3SD0089

Matrix: W-SURF-SLT

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1981579	Enterolert/QT	ENTEROCOCCI-ENTEROLERT	4840	J	MPN/100 mL		

Ref. Method and Comment:

Enterolert/QT: All Quanti Trays were positive. Results may be higher than reported. Lab did not properly dilute Marine Sample

Sample Location: North Creek(Tidal)@US41

Collection Date/Time: 04/04/2018 11:35

Field ID: G3SD0089_DUP

Matrix: W-SURF-SLT

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1981580	Enterolert/QT	ENTEROCOCCI-ENTEROLERT	2420	J	MPN/100 mL		

Ref. Method and Comment:

Enterolert/QT: All Quanti Trays were positive. Results may be higher than reported. Lab did not properly dilute Marine Sample

Sample Location: North Creek byOaksCrkCt

Collection Date/Time: 04/04/2018 12:00

Field ID: G3SD0064

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1981568	SM 10200 H (mod.)	Chlorophyll-a, Corrected	0.89	I	ug/L	P343380	
		Phaeophytin-a	0.85	I	ug/L	P343380	
		Chlorophyll-a, Uncorrected	1.4		ug/L	P343380	

Ref. Method and Comment:

SM 10200 H (mod.): Precision data is not available for at least one component due to the small amount of analyte in the QC sample. Refer to QA report for available precision data.

Sample Location: North Creek byOaksCrkCt

Collection Date/Time: 04/04/2018 12:00

Field ID: G3SD0064

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1981577	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray**	307.6		MPN/100 mL		

Sample Location: South Creek @ HonoreAve

Collection Date/Time: 04/04/2018 12:25

Field ID: G3SD0090

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1981578	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray**	12		MPN/100 mL		
1981667	SOP-PCR-4.0	HF183 - Human Source Marker**	Below Threshold				

Sample Location: North Creek(Tidal)@US41

Collection Date/Time: 04/04/2018 11:30

Field ID: G3SD0089

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1981665	SOP-PCR-1.0	HF183-qPCR**	130	T	GEU/100 mL	P343867	
	SOP-PCR-4.0	HF183 - Human Source Marker**	Above Threshold				

Sample Location: North Creek(Tidal)@US41

Collection Date/Time: 04/04/2018 11:35

Field ID: G3SD0089_DUP

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1981666	SOP-PCR-1.0	HF183-qPCR**	510	I	GEU/100 mL	P343867	
	SOP-PCR-4.0	HF183 - Human Source Marker**	Above Threshold				

Quality Assurance Report Method Blank Results

Reference Method: SM 10200 H (mod.)

Batch ID: P343380

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: P343867

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

Quality Assurance Report Laboratory Control Sample Accuracy

Reference Method: SM 10200 H (mod.)

Batch ID: P343380

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	85.5		P	84 - 116

Quality Assurance Report Laboratory Control Sample Accuracy

Reference Method: SOP-PCR-1.0
 Batch ID: P343867

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	95.7	70.1	P/P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
 Batch ID: P343867

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
1976854	HF183-qPCR	72.1		P	50 - 175
1976855	HF183-qPCR	61.9		P	50 - 175
1976857	HF183-qPCR	55.5		P	50 - 175
1976858	HF183-qPCR	56.2		P	50 - 175
1976859	HF183-qPCR	63.5		P	50 - 175
1980254	HF183-qPCR	74.2		P	50 - 175
1981665	HF183-qPCR	84.2		P	50 - 175
1981666	HF183-qPCR	76.2		P	50 - 175
1982100	HF183-qPCR	79.6		P	50 - 175
1983688	HF183-qPCR	77.3		P	50 - 175
1984031	HF183-qPCR	82.1		P	50 - 175
1987386	HF183-qPCR	80.4		P	50 - 175
1987512	HF183-qPCR	74.8	95.8	P/P	50 - 175

Quality Assurance Report Precision

Reference Method: SM 10200 H (mod.)
 Batch ID: P343380

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
1981485	Chlorophyll-a, Corrected	3.57	Sample	P	0 - 25
1981485	Chlorophyll-a, Uncorrected	4.50	Sample	P	0 - 25

Reference Method: SOP-PCR-1.0
 Batch ID: P343867

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
1987512	HF183-qPCR	24.6	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

Quality Assurance Report Surrogates

Lab Sample ID: 1981665
Field Sample ID: G3SD0089

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	110	P	50 - 300

Lab Sample ID: 1981666
Field Sample ID: G3SD0089_DUP

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	116	P	50 - 300

Quality Assurance Report Summary

Ref. Method	Analyte	LCS % Recovery	MS % Recovery	LCS	Precision SMP	MS
SM 10200 H (mod.)	Chlorophyll-a, Corrected	85.5			3.57	
	Chlorophyll-a, Uncorrected				4.50	
SOP-PCR-1.0	HF183-qPCR	70.1 95.7	76.2 74.2 80.4 79.6			24.6

Reference Method Descriptions

Method / DoH Cert #	Description	Associated Samples
Enterolert/QT / E85457	Enterococci by MPN method by Sanders Laboratories Overflow Laboratory for SD	1981579, 1981580
SM 10200 H (mod.) / E31780	Phytoplankton chlorophyll-a corrected, uncorrected, and phaeophytin by spectrophotometry	1981567, 1981568
SM 9223 Quanti-Tray / E85457	Escherichia coli by Colilert 18 Quanti-Tray method by Sanders Laboratories for South District	1981573, 1981574, 1981577, 1981578
SOP-PCR-1.0 / E31780	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	1981665, 1981666
SOP-PCR-4.0 / E31780	Preparation of Samples by Filtration and held for qPCR Analysis	1981664, 1981665, 1981666, 1981667

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
Enterolert/QT	04/04/2018	04/04/2018 14:40	Edwin Gonzalez-Soto	04/05/2018 14:59	Sanders-Fort Myers	1981579, 1981580
SM 10200 H (mod.)	04/05/2018	04/05/2018 14:45		04/12/2018 15:45	Jennifer Mihalic	1981567, 1981568
SM 9223 Quanti-Tray	04/04/2018	04/04/2018 15:22		04/05/2018 10:08	Sanders-Fort Myers	1981578
	04/04/2018	04/04/2018 15:33		04/05/2018 10:08	Sanders-Fort Myers	1981573, 1981574, 1981577
SOP-PCR-1.0	04/05/2018	04/05/2018 13:30	Puja Jasrotia	05/02/2018 17:03	Mailin Lopez	1981665, 1981666
SOP-PCR-4.0	04/05/2018	04/05/2018 13:30	Angela Smith	04/23/2018 10:15	Puja Jasrotia	1981664, 1981665, 1981666, 1981667