

Biological Analysis Report

CEN-DIST-2018-02-06-03

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: **Long Branch / Crane Strand / CS Drain / UnDrain**
Request ID: **RQ-2018-02-05-73**
Customer: **CEN-DIST**
Project ID: **SMP**

Send Reports to:
FL Dept. of Environmental Protection
FL Dept. of Environmental Protection
3319 Maguire Blvd., Suite 232
Orlando, FL 32803-3767
Attn: Mike Linger

For additional information please contact
Cheryl Swanson - Administrator
J. Marshall Faircloth - Aquatic Toxicology
Jacqueline Savage - Bench Biology
Jacqueline Savage - Microbiology
Puja Jasrotia, Ph.D. - Molecular Biology
Jennifer Walters - Taxonomy (Invertebrate & Algal)
Thekkekalathil Chandrasekhar, Ph.D, QA Officer
Phone (850) 245-8177

Certified by: Cheryl A. Swanson, Program Administrator

Date Certified: 13-MAR-2018 08:54



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: UNNAMED DRAIN @ LAKE LIVE OAK RD.

Collection Date/Time: 02/05/2018 11:42

Field ID: G2CE0168

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1964955	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray	196		MPN/100 mL		
1964961	SM 10200 H (mod.)	Chlorophyll-a, Corrected	2.0		ug/L	P339946	
		Phaeophytin-a	0.40	U	ug/L	P339946	
		Chlorophyll-a, Uncorrected	2.1		ug/L	P339946	

Sample Location: UNNAMED DRAIN @ AVALON PARK BLVD

Collection Date/Time: 02/05/2018 12:10

Field ID: G2CE0169

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1964956	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray	276		MPN/100 mL		
1964962	SM 10200 H (mod.)	Chlorophyll-a, Corrected	0.99	I	ug/L	P339946	
		Phaeophytin-a	0.40	U	ug/L	P339946	
		Chlorophyll-a, Uncorrected	1.1	I	ug/L	P339946	

Sample Location: CRANE STRAND @ 100M DS OF ECON TRL BRIDGE

Collection Date/Time: 02/05/2018 13:07

Field ID: G2CE0076

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1964957	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray	172		MPN/100 mL		

Sample Location: CRANE STRAND AT HERRELL RD.

Collection Date/Time: 02/05/2018 13:21

Field ID: G2CE0156

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1964952	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray	326		MPN/100 mL		

Sample Location: CRANE STRAND DRAIN @ 100M US OF UNIVERSITY BLVD

Collection Date/Time: 02/05/2018 13:52

Field ID: G2CE0077

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1964953	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray	1550		MPN/100 mL		
1964959	SM 10200 H (mod.)	Chlorophyll-a, Corrected	1.9		ug/L	P339941	
		Phaeophytin-a	0.50	I	ug/L	P339941	
		Chlorophyll-a, Uncorrected	2.3		ug/L	P339941	
1964973	SOP-PCR-1.0	HF183-qPCR**	6.47E+03		GEU/100 mL	P340649	
	SOP-PCR-4.0	HF183 - Human Source Marker**	Above Threshold				

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: CRANE STRAND DRAIN @ 250M NORTH OF UNIVERSITY BLVD

Collection Date/Time: 02/05/2018 14:18

Field ID: G2CE0157

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
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Field ID: G2CE0157

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1964954	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray	1120		MPN/100 mL		
1964960	SM 10200 H (mod.)	Chlorophyll-a, Corrected	0.97	I	ug/L	P339941	
		Phaeophytin-a	0.40	U	ug/L	P339941	
		Chlorophyll-a, Uncorrected	1.2	I	ug/L	P339941	
1964974	SOP-PCR-1.0	HF183-qPCR**	52	T	GEU/100 mL	P340649	
	SOP-PCR-4.0	HF183 - Human Source Marker**	Above Threshold				

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.

Quality Assurance Report
Method Blank Results

Reference Method: SM 10200 H (mod.)
Batch ID: P339941

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: SM 10200 H (mod.)
Batch ID: P339946

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: P340649

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: P339941

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	103		P	84 - 116

Reference Method: SM 10200 H (mod.)
Batch ID: P339946

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	104		P	84 - 116

Reference Method: SOP-PCR-1.0
Batch ID: P340649

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
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Quality Assurance Report Laboratory Control Sample Accuracy

Reference Method: SOP-PCR-1.0
 Batch ID: P340649

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	117	86.1	P/P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
 Batch ID: P340649

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
1964973	HF183-qPCR	74.0		P	50 - 175
1964974	HF183-qPCR	84.2		P	50 - 175
1965458	HF183-qPCR	91.9		P	50 - 175
1965459	HF183-qPCR	70.6		P	50 - 175
1965461	HF183-qPCR	83.3		P	50 - 175
1966060	HF183-qPCR	67.1		P	50 - 175
1966063	HF183-qPCR	68.5		P	50 - 175
1966064	HF183-qPCR	72.6		P	50 - 175
1966327	HF183-qPCR	93.3		P	50 - 175
1971016	HF183-qPCR	87.0	81.6	P/P	50 - 175

Quality Assurance Report Precision

Reference Method: SM 10200 H (mod.)
 Batch ID: P339941

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
1964848	Chlorophyll-a, Corrected	12.7	Sample	P	0 - 25
1964848	Chlorophyll-a, Uncorrected	13.8	Sample	P	0 - 25

Reference Method: SM 10200 H (mod.)
 Batch ID: P339946

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
1965239	Chlorophyll-a, Corrected	8.91	Sample	P	0 - 25
1965239	Chlorophyll-a, Uncorrected	5.89	Sample	P	0 - 25
1965239	Phaeophytin-a	0.544	Sample	P	0 - 25

Reference Method: SOP-PCR-1.0
 Batch ID: P340649

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
1971016	HF183-qPCR	6.39	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: 1964973
Field Sample ID: G2CE0077

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	102	P	50 - 300

Lab Sample ID: 1964974
Field Sample ID: G2CE0157

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	102	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	103			12.7	
	Chlorophyll-a, Corrected	104			8.91	
	Chlorophyll-a, Uncorrected				13.8	
	Chlorophyll-a, Uncorrected				5.89	
	Phaeophytin-a				0.544	
SOP-PCR-1.0	HF183-qPCR	86.1 117	91.9 70.6 83.3 67.1			6.39

Reference Method Descriptions

Method / DoH Cert #	Description	Associated Samples
SM 10200 H (mod.) / E31780	Phytoplankton chlorophyll-a corrected, uncorrected, and phaeophytin by spectrophotometry	1964959, 1964960, 1964961, 1964962
SM 9223 Quanti-Tray / E83018	Escherichia coli by Colilert 18 Quanti-Tray method by FlowersOverflow Laboratory for Central District	1964952, 1964953, 1964954, 1964955, 1964956, 1964957
SOP-PCR-1.0 / E31780	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	1964973, 1964974
SOP-PCR-4.0 / E31780	Preparation of Samples by Filtration and held for qPCR Analysis	1964973, 1964974

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
SM 10200 H (mod.)	02/06/2018	02/06/2018 14:11	Cathy Sherrer	02/13/2018	Jennifer Mihalic	1964959, 1964960, 1964961, 1964962
SM 9223 Quanti-Tray	02/05/2018	02/05/2018 15:00		02/06/2018 10:15	Flowers - Altamonte Springs	1964952, 1964953, 1964954, 1964955, 1964956, 1964957
SOP-PCR-1.0	02/06/2018	02/06/2018 14:15	Puja Jasrotia	03/08/2018	Sarah Menz	1964973, 1964974
SOP-PCR-4.0	02/06/2018	02/06/2018 14:15	Angela Smith	03/07/2018	Puja Jasrotia	1964973, 1964974