

Biological Analysis Report

WAS-SECT-2018-03-22-01

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

Event Description: **CDD, Godby, Megginnis, Chicken, Sand, 8, Black158, Big Branch**
Request ID: **RQ-2018-03-19-26**
Customer: **WAS-SECT**
Project ID: **SMP**

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Date Certified: 24-APR-2018 13:20



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, Microbiology and 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 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: SAND POND MIDDLE

Collection Date/Time: 03/22/2018 11:15

Field ID: G1WA0044

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1976987	SM 10200 H (mod.)	Chlorophyll-a, Corrected	2.9		ug/L	P342317	
		Phaeophytin-a	2.0		ug/L	P342317	
		Chlorophyll-a, Uncorrected	4.2		ug/L	P342317	

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: SAND POND MIDDLE

Collection Date/Time: 03/22/2018 11:15

Field ID: G1WA0044

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1976993	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray	8.0	A	MPN/100 mL	P342239	

Sample Location: EIGHT MILE POND MIDDLE

Collection Date/Time: 03/22/2018 11:50

Field ID: G1WA0024

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1976988	SM 10200 H (mod.)	Chlorophyll-a, Corrected	40		ug/L	P342317	
		Phaeophytin-a	1.8		ug/L	P342317	
		Chlorophyll-a, Uncorrected	42		ug/L	P342317	

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: EIGHT MILE POND MIDDLE

Collection Date/Time: 03/22/2018 11:50

Field ID: G1WA0024

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1976994	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray	131.4		MPN/100 mL	P342239	

Sample Location: BLACK CREEK UPSTREAM OF CR 158

Collection Date/Time: 03/22/2018 13:05

Field ID: G1WA0072

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1976989	SM 10200 H (mod.)	Chlorophyll-a, Corrected	1.7		ug/L	P342317	
		Phaeophytin-a	0.78	I	ug/L	P342317	
		Chlorophyll-a, Uncorrected	2.3		ug/L	P342317	

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: BLACK CREEK UPSTREAM OF CR 158

Collection Date/Time: 03/22/2018 13:05

Field ID: G1WA0072

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1976995	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray	161.6		MPN/100 mL	P342239	

Sample Location: BLACK CREEK UPSTREAM OF CR 158

Collection Date/Time: 03/22/2018 13:05

Field ID: G1WA0072

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1977001	SOP-PCR-4.0	HF183 - Human Source Marker**	Below Threshold				

Sample Location: CENTRAL DRAINAGE DITCH DOWNSTREAM OF ORANGE AVE Collection Date/Time: 03/22/2018 10:00

Field ID: G1WA0026

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1976990	SM 10200 H (mod.)	Chlorophyll-a, Corrected	16		ug/L	P342317	
		Phaeophytin-a	3.1		ug/L	P342317	
		Chlorophyll-a, Uncorrected	19		ug/L	P342317	

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: CENTRAL DRAINAGE DITCH DOWNSTREAM OF ORANGE AVE Collection Date/Time: 03/22/2018 10:00

Field ID: G1WA0026

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1976996	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray	378.4		MPN/100 mL	P342239	

Sample Location: CHICKEN BRANCH AT OLD PLANK RD Collection Date/Time: 03/22/2018 12:31

Field ID: G1WA0031

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1976997	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray	195.6		MPN/100 mL	P342239	

Sample Location: CHICKEN BRANCH AT OLD PLANK RD Collection Date/Time: 03/22/2018 12:31

Field ID: G1WA0031

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1977002	SOP-PCR-4.0	HF183 - Human Source Marker**	Below Threshold				

Sample Location: BIG BRANCH 50 METERS UPSTREAM OF CR 67 Collection Date/Time: 03/22/2018 08:25

Field ID: G1WA0001

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1976998	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray	127.4		MPN/100 mL	P342239	

Sample Location: BIG BRANCH 50 METERS UPSTREAM OF CR 67 Collection Date/Time: 03/22/2018 08:25

Field ID: G1WA0001

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1977003	SOP-PCR-4.0	HF183 - Human Source Marker**	Below Threshold				

Sample Location: GODBY DRAINAGE AT PROGRESS DR Collection Date/Time: 03/22/2018 09:25

Field ID: G1WA0061

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1976991	SM 10200 H (mod.)	Chlorophyll-a, Corrected	0.58	I	ug/L	P342317	
		Phaeophytin-a	0.51	I	ug/L	P342317	
		Chlorophyll-a, Uncorrected	0.91	I	ug/L	P342317	

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: GODBY DRAINAGE AT PROGRESS DR

Collection Date/Time: 03/22/2018 09:25

Field ID: G1WA0061

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1976999	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray	686.7		MPN/100 mL	P342239	

Sample Location: MEGGINNIS ARM RUN DOWNSTREAM OF SHARER

Collection Date/Time: 03/22/2018 07:25

Field ID: G1WA0033

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1976992	SM 10200 H (mod.)	Chlorophyll-a, Corrected	0.94	I	ug/L	P342317	
		Phaeophytin-a	1.0	I	ug/L	P342317	
		Chlorophyll-a, Uncorrected	1.6		ug/L	P342317	

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: MEGGINNIS ARM RUN DOWNSTREAM OF SHARER

Collection Date/Time: 03/22/2018 07:25

Field ID: G1WA0033

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1977000	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray	369.0		MPN/100 mL	P342239	

Sample Location: MEGGINNIS ARM RUN DOWNSTREAM OF SHARER

Collection Date/Time: 03/22/2018 07:25

Field ID: G1WA0033

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1977004	SOP-PCR-4.0	HF183 - Human Source Marker**	Below Threshold				

Quality Assurance Report Method Blank Results

Reference Method: SM 10200 H (mod.)

Batch ID: P342317

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

Quality Assurance Report Method Blank Results

Reference Method: SM 9223 Quanti-Tray
Batch ID: P342239

Component	Result	Code	Units
Escherichia Coli-Quanti-Tray	1.0	U	MPN/100 mL

Quality Assurance Report Laboratory Control Sample Accuracy

Reference Method: SM 10200 H (mod.)
Batch ID: P342317

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	98.9		P	84 - 116

Quality Assurance Report Precision

Reference Method: SM 10200 H (mod.)
Batch ID: P342317

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
1977087	Chlorophyll-a, Corrected	3.67	Sample	P	0 - 25
1977087	Chlorophyll-a, Uncorrected	1.28	Sample	P	0 - 25

Reference Method: SM 9223 Quanti-Tray
Batch ID: P342239

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
1976713	Escherichia Coli-Quanti-Tray	17.4	Sample	P	0 - 80
1976993	Escherichia Coli-Quanti-Tray	70.8	Sample	P	0 - 80

* 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 Summary

Ref. Method	Analyte	LCS % Recovery	MS % Recovery	LCS	Precision SMP	MS
SM 10200 H (mod.)	Chlorophyll-a, Corrected	98.9			3.67	
	Chlorophyll-a, Uncorrected				1.28	
SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray				17.4	70.8

Reference Method Descriptions

Method / DoH Cert #	Description	Associated Samples
SM 10200 H (mod.) / E31780	Phytoplankton chlorophyll-a corrected, uncorrected, and phaeophytin by spectrophotometry	1976987, 1976988, 1976989, 1976990, 1976991, 1976992
SM 9223 Quanti-Tray / E31780	Escherichia coli by Colilert 18 Quanti-Tray method - non-chlorinated water systems	1976993, 1976994, 1976995, 1976996, 1976997, 1976998, 1976999, 1977000
SOP-PCR-4.0 / E31780	Preparation of Samples by Filtration and held for qPCR Analysis	1977001, 1977002, 1977003, 1977004

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
SM 10200 H (mod.)	03/22/2018	03/22/2018 15:59	Cathy Sherrer	03/27/2018 13:05	Kim M. Spencer	1976987, 1976988, 1976989, 1976990, 1976991, 1976992
SM 9223 Quanti-Tray	03/22/2018	03/22/2018 15:20	Edwin Gonzalez-Soto	03/23/2018 09:55	Edwin Gonzalez-Soto	1976993, 1976994, 1976995, 1976996, 1976997, 1976998, 1976999, 1977000
SOP-PCR-4.0	03/22/2018	03/22/2018 15:00	Sarah Menz	04/18/2018 15:52	Puja Jasrotia	1977001, 1977002, 1977003, 1977004