

Biological Analysis Report

NE-DIST-2017-02-01-01

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. - North
812 S.W. Harvey Greene Drive
Madison, Florida 32341
DOH Accreditation E82405

Event Description: **Suwannee North 2**

Request ID: **RQ-2017-01-30-10**

Customer: **NE-DIST**

Project ID: **SMP**

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8800 Baymeadows Way West
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Certified by: Cheryl A. Swanson, Environmental Manager

Date Certified: 07-MAR-2017 14:35



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: Falling Creek @ US 441

Collection Date/Time: 01/31/2017 09:20

Field ID: G1NE0236 01312017

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1868533	SM 10200 H (mod.)	Chlorophyll-a, Corrected	0.55	U	ug/L	P319376	
		Phaeophytin-a	0.40	U	ug/L	P319376	
		Chlorophyll/Phaeophytin Ratio	0.98			P319376	
		Chlorophyll-a, Uncorrected	0.40	U	ug/L	P319376	
1868560	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray	225		MPN/100 mL		

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: Falling Creek @ US 441

Collection Date/Time: 01/31/2017 09:20

Field ID: G1NE0236 01312017

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1868572	SOP-PCR-4.0	HF183 - Human Source Marker**	Below Threshold				

Sample Location: Deep Creek @ US 441

Collection Date/Time: 01/31/2017 09:55

Field ID: G1NE0267 01312017

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1868534	SM 10200 H (mod.)	Chlorophyll-a, Corrected	0.55	U	ug/L	P319376	
		Phaeophytin-a	0.40	U	ug/L	P319376	
		Chlorophyll/Phaeophytin Ratio	2.0			P319376	
		Chlorophyll-a, Uncorrected	0.40	U	ug/L	P319376	
1868561	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray	60.2		MPN/100 mL		

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: Deep Creek @ US 441

Collection Date/Time: 01/31/2017 09:55

Field ID: G1NE0267 01312017

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1868573	SOP-PCR-4.0	HF183 - Human Source Marker**	Below Threshold				

Sample Location: Little Creek @ US 441

Collection Date/Time: 01/31/2017 10:20

Field ID: G1NE0251 01312017

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1868535	SM 10200 H (mod.)	Chlorophyll-a, Corrected	0.55	U	ug/L	P319377	
		Phaeophytin-a	0.40	U	ug/L	P319377	
		Chlorophyll/Phaeophytin Ratio	1.7			P319377	
		Chlorophyll-a, Uncorrected	0.40	U	ug/L	P319377	
1868562	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray	613		MPN/100 mL		

Sample Location: Little Creek @ US 441

Collection Date/Time: 01/31/2017 10:20

Field ID: G1NE0251 01312017

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1868574	SOP-PCR-1.0	HF183-qPCR**	130	I	GEU/100 mL	P319590	
	SOP-PCR-4.0	HF183 - Human Source Marker**	Above Threshold				

Sample Location: Hunters Creek @ SR 135

Collection Date/Time: 01/31/2017 10:55

Field ID: G1NE0384 01312017

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1868566	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray	1120		MPN/100 mL		
1868575	SOP-PCR-1.0	HF183-qPCR**	35	T	GEU/100 mL	P319590	
	SOP-PCR-4.0	HF183 - Human Source Marker**	Above Threshold				

Sample Location: Swift Creek @ CR 25A

Collection Date/Time: 01/31/2017 11:30

Field ID: G1NE0327 01312017

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1868567	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray	119		MPN/100 mL		
1868576	SOP-PCR-4.0	HF183 - Human Source Marker**	Below Threshold				

Sample Location: Camp Branch @ CR 25A

Collection Date/Time: 01/31/2017 11:50

Field ID: G1NE0269 01312017

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1868568	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray	194		MPN/100 mL		
1868577	SOP-PCR-4.0	HF183 - Human Source Marker**	Below Threshold				

Quality Assurance Report Method Blank Results

Reference Method: SM 10200 H (mod.)
Batch ID: P319376

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: P319377

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: P319590

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: P319376

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	92.9		P	84 - 116

Reference Method: SM 10200 H (mod.)
Batch ID: P319377

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	87.1		P	84 - 116

Reference Method: SOP-PCR-1.0
Batch ID: P319590

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	93.3	76.8	P/P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
Batch ID: P319590

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
1864645	HF183-qPCR	74.4		P	50 - 175
1866758	HF183-qPCR	89.3		P	50 - 175
1866855	HF183-qPCR	143		P	50 - 175
1868574	HF183-qPCR	132		P	50 - 175
1868575	HF183-qPCR	116		P	50 - 175
1868829	HF183-qPCR	110		P	50 - 175
1871159	HF183-qPCR	72.5		P	50 - 175
1871160	HF183-qPCR	108	136	P/P	50 - 175
1871161	HF183-qPCR	132		P	50 - 175
1871162	HF183-qPCR	85.8		P	50 - 175
1871755	HF183-qPCR	117		P	50 - 175
1871756	HF183-qPCR	87.9		P	50 - 175

Quality Assurance Report Precision

Reference Method: SM 10200 H (mod.)
Batch ID: P319376

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
1868488	Chlorophyll/Phaeophytin Ratio	0.0	Sample	P	0 - 25
1868488	Chlorophyll-a, Corrected	3.77	Sample	P	0 - 25
1868488	Chlorophyll-a, Uncorrected	2.96	Sample	P	0 - 25

Reference Method: SM 10200 H (mod.)
Batch ID: P319377

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
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Quality Assurance Report Precision

Reference Method: SM 10200 H (mod.)
Batch ID: P319377

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
1868617	Chlorophyll/Phaeophytin Ratio	0.0	Sample	P	0 - 25
1868617	Chlorophyll-a, Corrected	4.99	Sample	P	0 - 25
1868617	Chlorophyll-a, Uncorrected	5.94	Sample	P	0 - 25
1868617	Phaeophytin-a	10.9	Sample	P	0 - 25

Reference Method: SOP-PCR-1.0
Batch ID: P319590

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
1871160	HF183-qPCR	6.13	Sample	P	0 - 25
1871160	HF183-qPCR	22.6	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

Lab Sample ID: 1868574
Field Sample ID: G1NE0251 01312017

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	105	P	50 - 300

Lab Sample ID: 1868575
Field Sample ID: G1NE0384 01312017

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	106	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/Phaeophytin Ratio							0.0	
	Chlorophyll-a, Corrected	92.9						3.77	
	Chlorophyll-a, Corrected	87.1						4.99	
	Chlorophyll-a, Uncorrected							2.96	
	Chlorophyll-a, Uncorrected							5.94	
	Phaeophytin-a							10.9	
SOP-PCR-1.0	HF183-qPCR	76.8	93.3	87.9	72.5	108		6.13	22.6
				136					

Reference Method Descriptions

Method / DoH Cert #	Description	Associated Samples
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Reference Method Descriptions

Method / DoH Cert #	Description	Associated Samples
SM 10200 H (mod.) / E31780	Phytoplankton chlorophyll-a corrected, uncorrected, phaeophytin and ratio by spectrophotometry	1868533, 1868534, 1868535
SM 9223 Quanti-Tray / E82405	Escherichia coli by Colilert 18 Quanti-Tray method by FlowersOverflow Laboratory for Northeast District	1868560, 1868561, 1868562, 1868566, 1868567, 1868568
SOP-PCR-1.0 / E31780	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	1868574, 1868575
SOP-PCR-4.0 / E31780	Preparation of Samples by Filtration and held for qPCR Analysis	1868572, 1868573, 1868574, 1868575, 1868576, 1868577

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
SM 10200 H (mod.)	02/01/2017	02/01/2017 13:40	Morgan Lane	02/06/2017	Brian Cumbie	1868533, 1868534, 1868535
SM 9223 Quanti-Tray	01/31/2017			01/31/2017 14:27	Flower-MAD	1868560, 1868561, 1868562, 1868566, 1868567, 1868568
SOP-PCR-1.0	02/01/2017	02/01/2017 13:00	Sarah Menz	02/15/2017	Sarah Menz	1868574, 1868575
SOP-PCR-4.0	02/01/2017	02/01/2017 13:00	Avril Wood	02/14/2017	Daisys Matthews	1868572, 1868573, 1868574, 1868575, 1868576, 1868577