

Biological Analysis Report

SE-DIST-2017-10-10-01

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.
571 NW Mercantile Pl. Ste. 111
Port St. Lucie, FL 34986
DOH Accreditation E86562

Event Description: **SMP-G3 WPB Run**
Request ID: **RQ-2017-10-09-29**
Customer: **SE-DIST**
Project ID: **SMP**

Send Reports to:
FL Dept. of Environmental Protection
FL Dept. of Environmental Protection
2199 South Rock Road
Fort Pierce, FL 34945
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Date Certified: 28-NOV-2017 17:29



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: Acme South @ Lake Worth Rd.

Collection Date/Time: 10/09/2017 10:05

Field ID: G3SE0016

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1935237	SM 10200 H (mod.)	Chlorophyll-a, Corrected	15		ug/L	P333374	
		Phaeophytin-a	5.2		ug/L	P333374	
		Chlorophyll-a, Uncorrected	19		ug/L	P333374	

Sample Location: Acme North @ South Shore Blvd

Collection Date/Time: 10/09/2017 10:25

Field ID: 32012006

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1935238	SM 10200 H (mod.)	Chlorophyll-a, Corrected	15		ug/L	P333374	
		Phaeophytin-a	4.3		ug/L	P333374	
		Chlorophyll-a, Uncorrected	18		ug/L	P333374	

Sample Location: Okeeheele Park Boat Ramp

Collection Date/Time: 10/09/2017 11:00

Field ID: G3SE0015

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1935243	SM 10200 H (mod.)	Chlorophyll-a, Corrected	3.7		ug/L	P333374	
		Phaeophytin-a	0.42	I	ug/L	P333374	
		Chlorophyll-a, Uncorrected	4.1		ug/L	P333374	

Sample Location: Lake Clarke @ Pine Tree Lane

Collection Date/Time: 10/09/2017 11:30

Field ID: 28010809

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1935242	SM 10200 H (mod.)	Chlorophyll-a, Corrected	21		ug/L	P333374	
		Phaeophytin-a	3.2		ug/L	P333374	
		Chlorophyll-a, Uncorrected	24		ug/L	P333374	
1935259	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray	85.0		MPN/100 mL		

Sample Location: Pine Lake

Collection Date/Time: 10/09/2017 12:00

Field ID: 32008010

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1935241	SM 10200 H (mod.)	Chlorophyll-a, Corrected	22		ug/L	P333374	
		Phaeophytin-a	6.2		ug/L	P333374	
		Chlorophyll-a, Uncorrected	26		ug/L	P333374	
1935252	SOP-PCR-1.0	HF183-qPCR**	12	T	GEU/100 mL	P334675	
	SOP-PCR-4.0	HF183 - Human Source Marker**	Above Threshold				
1935258	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray	441		MPN/100 mL		

Sample Location: Clear Lake @ Okeechobee Rd.

Collection Date/Time: 10/09/2017 12:20

Field ID: 28010896

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1935239	SM 10200 H (mod.)	Chlorophyll-a, Corrected	9.8		ug/L	P333374	

Field ID: 28010896

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1935239	SM 10200 H (mod.)	Phaeophytin-a	0.70	I	ug/L	P333374	
		Chlorophyll-a, Uncorrected	11		ug/L	P333374	

Sample Location: C-17 @ SR-702

Collection Date/Time: 10/09/2017 13:00

Field ID: 32008003

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1935240	SM 10200 H (mod.)	Chlorophyll-a, Corrected	31		ug/L	P333374	
		Phaeophytin-a	5.9		ug/L	P333374	
		Chlorophyll-a, Uncorrected	36		ug/L	P333374	
1935257	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray	246		MPN/100 mL		

Quality Assurance Report Method Blank Results

Reference Method: SM 10200 H (mod.)
 Batch ID: P333374

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: P334675

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

Quality Assurance Report Laboratory Control Sample Accuracy

Reference Method: SM 10200 H (mod.)
 Batch ID: P333374

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

Reference Method: SOP-PCR-1.0
 Batch ID: P334675

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	82.6	107	P/P	50 - 175
HF183-qPCR	81.7	95.7	P/P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
Batch ID: P334675

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
1935252	HF183-qPCR	79.2		P	50 - 175
1936509	HF183-qPCR	93.5		P	50 - 175
1936510	HF183-qPCR	93.3		P	50 - 175
1936867	HF183-qPCR	73.4		P	50 - 175
1936920	HF183-qPCR	61.3		P	50 - 175
1937848	HF183-qPCR	75.7		P	50 - 175
1942953	HF183-qPCR	91.2	81.5	P/P	50 - 175
1943020	HF183-qPCR	92.1		P	50 - 175
1944326	HF183-qPCR	79.1		P	50 - 175
1944327	HF183-qPCR	76.5		P	50 - 175
1944328	HF183-qPCR	79.3		P	50 - 175
1944330	HF183-qPCR	90.1		P	50 - 175

Quality Assurance Report Precision

Reference Method: SM 10200 H (mod.)
Batch ID: P333374

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
1935429	Chlorophyll-a, Corrected	11.4	Sample	P	0 - 25
1935429	Chlorophyll-a, Uncorrected	8.71	Sample	P	0 - 25
1935429	Phaeophytin-a	14.0	Sample	P	0 - 25

Reference Method: SOP-PCR-1.0
Batch ID: P334675

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
1942953	HF183-qPCR	11.2	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: 1935252
Field Sample ID: 32008010

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	87.0	P	50 - 300

Quality Assurance Report Summary

Ref. Method	Analyte	LCS % Recovery	MS % Recovery	Precision
			LCS	SMP
				MS
SM 10200 H (mod.)	Chlorophyll-a, Corrected	104		11.4

Quality Assurance Report Summary

Ref. Method	Analyte	LCS % Recovery			MS % Recovery			LCS	Precision	MS
									SMP	
SM 10200 H (mod.)	Chlorophyll-a, Uncorrected								8.71	
	Phaeophytin-a								14.0	
SOP-PCR-1.0	HF183-qPCR	82.6	95.7	81.7	61.3	91.2	81.5			11.2
		107			92.1					

Reference Method Descriptions

Method / DoH Cert #	Description	Associated Samples
SM 10200 H (mod.) / E31780	Phytoplankton chlorophyll-a corrected, uncorrected, and phaeophytin by spectrophotometry	1935237, 1935238, 1935239, 1935240, 1935241, 1935242, 1935243
SM 9223 Quanti-Tray / E86562	Escherichia coli by Colilert 18 Quanti-Tray method by FlowersOverflow Laboratory for Southeast District	1935257, 1935258, 1935259
SOP-PCR-1.0 / E31780	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	1935252
SOP-PCR-4.0 / E31780	Preparation of Samples by Filtration and held for qPCR Analysis	1935252

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
SM 10200 H (mod.)	10/10/2017	10/10/2017 12:30	Cathy Sherrer	10/12/2017	Kim M. Spencer	1935237, 1935238, 1935239, 1935240, 1935241, 1935242, 1935243
SM 9223 Quanti-Tray	10/09/2017	10/09/2017 16:20		10/10/2017 11:50	Flowers-PSL	1935257, 1935258, 1935259
SOP-PCR-1.0	10/10/2017	10/10/2017 14:30	Avril Wood	11/15/2017	Sarah Menz	1935252
SOP-PCR-4.0	10/10/2017	10/10/2017 14:30	Avril Wood	11/08/2017	Puja Jasrotia	1935252