

Biological Analysis Report

SO-DIST-2019-08-07-01

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

Overflow Analyses Performed By:
Benchmark EnviroAnalytical, Inc. - North Port
1001 Corporate Avenue, Suite 102
North Port, FL 34289
DOH Accreditation E85086

Event Description: **2019 SMP : MyBB**

Request ID: **RQ-2019-08-05-41**

Customer: **SO-DIST**

Project ID: **SMP**

Send Reports to:
FL Dept. of Environmental Protection
FLDEP South District Office
2295 Victoria Ave. Suite 364
Fort Myers, FL 33901-3381
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Certified by: Puja Jasrotia, Environmental Manager

Date Certified: 11-SEP-2019 11:50



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 by the Florida Department of Health Environmental Laboratory Certification Program 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: LakeMyakka(Upper)

Collection Date/Time: 08/06/2019 10:50

Field ID: G3SD0133

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2109149	SM 10200 H (mod.)	Chlorophyll-a, Corrected	7.0		ug/L	P368961	
		Phaeophytin-a	1.7	I	ug/L	P368961	
		Chlorophyll-a, Uncorrected	8.2		ug/L	P368961	

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: Brandy Branch @SR70

Collection Date/Time: 08/06/2019 12:45

Field ID: G3SD0083

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2109148	SM 10200 H (mod.)	Chlorophyll-a, Corrected	1.8	I	ug/L	P368961	
		Phaeophytin-a	1.0	U	ug/L	P368961	
		Chlorophyll-a, Uncorrected	2.2		ug/L	P368961	
2109152	SOP-PCR-1.0	HF183-qPCR**	280	UJ	GEU/100 mL	P368498	MS
	SOP-PCR-1.4	BacR-qPCR**	3.40E+05		TSC/100 mL	P368498	
2109153	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray	657		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.

SOP-PCR-1.0: Quality control failure(s) observed. See QA report for details.

Quality Assurance Report Method Blank Results

Reference Method: SM 10200 H (mod.)

Batch ID: P368961

Component	Result	Code	Units
Chlorophyll-a, Corrected	0.82	U	ug/L
Chlorophyll-a, Uncorrected	0.60	U	ug/L
Phaeophytin-a	0.90	U	ug/L

Reference Method: SOP-PCR-1.0

Batch ID: P368498

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

Reference Method: SOP-PCR-1.4

Batch ID: P368498

Component	Result	Code	Units
BacR-qPCR	3.0E+03	U	TSC/100 mL

Quality Assurance Report Laboratory Control Sample Accuracy

Reference Method: SM 10200 H (mod.)

Batch ID: P368961

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

Reference Method: SM 10200 H (mod.)
Batch ID: P368961

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	90.0		P	84 - 116

Reference Method: SOP-PCR-1.0
Batch ID: P368498

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	88.3	99.4	P/P	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P368498

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
BacR-qPCR	92.2	78.3	P/P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
Batch ID: P368498

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2109108	HF183-qPCR	70.3	79.4	P/P	50 - 175
2109152	HF183-qPCR	49.6		F	50 - 175
2109426	HF183-qPCR	81.8		P	50 - 175
2109427	HF183-qPCR	89.2		P	50 - 175
2109428	HF183-qPCR	64.7		P	50 - 175
2109620	HF183-qPCR	60.5		P	50 - 175
2109621	HF183-qPCR	73.7		P	50 - 175
2109622	HF183-qPCR	93.7		P	50 - 175
2109623	HF183-qPCR	80.6		P	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P368498

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2109108	BacR-qPCR	77.0	75.6	P/P	50 - 175
2109152	BacR-qPCR	57.9		P	50 - 175
2109426	BacR-qPCR	80.5		P	50 - 175
2109427	BacR-qPCR	80.3		P	50 - 175
2109428	BacR-qPCR	59.5		P	50 - 175
2109620	BacR-qPCR	53.4		P	50 - 175
2109621	BacR-qPCR	64.0		P	50 - 175
2109622	BacR-qPCR	83.1		P	50 - 175
2109623	BacR-qPCR	68.7		P	50 - 175

Quality Assurance Report Precision

Quality Assurance Report Precision

Reference Method: SM 10200 H (mod.)
Batch ID: P368961

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2109092	Chlorophyll-a, Corrected	7.34	Sample	P	0 - 25
2109092	Chlorophyll-a, Uncorrected	9.59	Sample	P	0 - 25

Reference Method: SOP-PCR-1.0
Batch ID: P368498

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2109108	HF183-qPCR	12.1	Spike	P	0 - 25

Reference Method: SOP-PCR-1.4
Batch ID: P368498

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2109108	BacR-qPCR	1.84	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: 2109152
Field Sample ID: G3SD0083

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	86.5	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	126	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	90.0					7.34		
	Chlorophyll-a, Uncorrected						9.59		
SOP-PCR-1.0	HF183-qPCR	99.4	88.3	70.3	79.4	49.6		12.1	
SOP-PCR-1.4	BacR-qPCR	78.3	92.2	81.8					
				77.0	75.6	57.9		1.84	
				80.5					

Reference Method Descriptions

Method / DoH Cert #	Description	Associated Samples
SM 10200 H (mod.) / E31780	Phytoplankton chlorophyll-a corrected, uncorrected, and phaeophytin by spectrophotometry	2109148, 2109149
SM 9223 Quanti-Tray / E85086	E. coli by Colilert/QT by BEA South in North Port	2109153

Reference Method Descriptions

Method / DoH Cert #	Description	<u>Associated Samples</u>
SOP-PCR-1.0 / E31780	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	2109152
SOP-PCR-1.4 / E31780	Detection and quantification of Ruminant specific Bacteroidetes BacR genetic marker with quantitative polymerase chain reaction	2109152

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
SM 10200 H (mod.)	08/07/2019	08/07/2019 12:00	Peterson Janvier	08/13/2019 11:08	Yuliya Danyuk	2109148, 2109149
SM 9223 Quanti-Tray	08/06/2019	08/06/2019 15:35		08/07/2019 16:12	Benchmark EnviroAnalytical in North Port	2109153
SOP-PCR-1.0	08/07/2019	08/07/2019 16:00	Sarah Menz	08/20/2019 08:20	Mailin Lopez	2109152
SOP-PCR-1.4	08/07/2019	08/07/2019 16:00	Sarah Menz	08/21/2019 12:09	Mailin Lopez	2109152