

Biological Analysis Report

WQAP-2018-09-28-01

Florida Department of Environmental Protection
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DOH Accreditation E31780

Overflow Analyses Performed By:
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Event Description: **Continuous Monitoring 2018**
Request ID: **RQ-2018-09-24-58**
Customer: **WQAP**
Project ID: **SMP**

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Date Certified: 15-NOV-2018 11:18



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: Craig Cr @ Hendricks Ave

Collection Date/Time: 09/27/2018 10:15

Field ID: 20030793

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2029546	SM 10200 H (mod.)	Chlorophyll-a, Corrected	9.3		ug/L	P353200	RPD
		Phaeophytin-a	0.60	U	ug/L	P353200	
		Chlorophyll-a, Uncorrected	9.8		ug/L	P353200	RPD

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: Craig Cr @ Hendricks Ave

Collection Date/Time: 09/27/2018 10:15

Field ID: 20030793

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2029548	SOP-PCR-1.0	HF183-qPCR**	140	U	GEU/100 mL	P352839	
	SOP-PCR-1.2	GULL2-qPCR**	8.4E+03	U	TSC/100 mL	P352839	
	SOP-PCR-1.3	GFD-purified-qPCR**	170	U	TSC/100 mL	P353626	LCS, MS
	SOP-PCR-1.4	BacR-qPCR**	1.5E+03	U	TSC/100 mL	P352839	LCS
	SOP-PCR-4.0	HF183 - Human Source Marker**	Above Threshold				

Ref. Method and Comment:

SOP-PCR-1.2: Precision data unavailable for Analysis batch A87385

Sample Location: Craig Cr @ Hendricks Ave

Collection Date/Time: 09/27/2018 10:15

Field ID: 20030793

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2029547	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	1850		MPN/100 mL		

Quality Assurance Report Method Blank Results

Reference Method: SM 10200 H (mod.)

Batch ID: P353200

Component	Result	Code	Units
Chlorophyll-a, Corrected	0.69	U	ug/L
Chlorophyll-a, Uncorrected	0.50	U	ug/L
Phaeophytin-a	0.75	U	ug/L

Reference Method: SOP-PCR-1.0

Batch ID: P352839

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

Reference Method: SOP-PCR-1.2

Batch ID: P352839

Component	Result	Code	Units
GULL2-qPCR	1.7E+04	U	TSC/100 mL
GULL2-qPCR	1.7E+04	U	TSC/100 mL
GULL2-qPCR	1.7E+04	U	TSC/100 mL

Quality Assurance Report Method Blank Results

Reference Method: SOP-PCR-1.3
Batch ID: P353626

Component	Result	Code	Units
GFD-purified-qPCR	340	U	TSC/100 mL

Reference Method: SOP-PCR-1.4
Batch ID: P352839

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

Quality Assurance Report Laboratory Control Sample Accuracy

Reference Method: SM 10200 H (mod.)
Batch ID: P353200

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	102		P	84 - 116

Reference Method: SOP-PCR-1.0
Batch ID: P352839

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	110	130	P/P	50 - 175

Reference Method: SOP-PCR-1.2
Batch ID: P352839

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
GULL2-qPCR	179	131	P/P	50 - 300
GULL2-qPCR	131	102	P/P	50 - 300
GULL2-qPCR	179	123	P/P	50 - 300

Reference Method: SOP-PCR-1.3
Batch ID: P353626

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
GFD-purified-qPCR	148	13.2	P/F	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P352839

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
BacR-qPCR	184	126	F/P	50 - 175
BacR-qPCR	311	102	F/P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0

Batch ID: P352839

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2028729	HF183-qPCR	100		P	50 - 175
2029548	HF183-qPCR	103		P	50 - 175
2032519	HF183-qPCR	95.7		P	50 - 175
2032845	HF183-qPCR	99.4		P	50 - 175
2032846	HF183-qPCR	102	96.0	P/P	50 - 175
2032847	HF183-qPCR	79.5		P	50 - 175
2032848	HF183-qPCR	84.4		P	50 - 175
2032849	HF183-qPCR	104		P	50 - 175
2033938	HF183-qPCR	99.7		P	50 - 175
2033939	HF183-qPCR	108		P	50 - 175
2034669	HF183-qPCR	103		P	50 - 175
2034670	HF183-qPCR	95.3		P	50 - 175
2034671	HF183-qPCR	100		P	50 - 175

Reference Method: SOP-PCR-1.2

Batch ID: P352839

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2029548	GULL2-qPCR	84.1	84.1	P/P	50 - 300
2032849	GULL2-qPCR	121		P	50 - 300

Reference Method: SOP-PCR-1.3

Batch ID: P353626

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2026896	GFD-purified-qPCR	0.0		F	50 - 175
2027411	GFD-purified-qPCR	114		P	50 - 175
2029548	GFD-purified-qPCR	107		P	50 - 175
2030524	GFD-purified-qPCR	114	134	P/P	50 - 175

Reference Method: SOP-PCR-1.4

Batch ID: P352839

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2029548	BacR-qPCR	88.1		P	50 - 175
2032519	BacR-qPCR	108		P	50 - 175
2032849	BacR-qPCR	91.7		P	50 - 175

Quality Assurance Report Precision

Reference Method: SM 10200 H (mod.)

Batch ID: P353200

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2029538	Chlorophyll-a, Corrected	25.5	Sample	F	0 - 25
2029538	Chlorophyll-a, Uncorrected	25.5	Sample	F	0 - 25

Quality Assurance Report Precision

Reference Method: SOP-PCR-1.0
Batch ID: P352839

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2032846	HF183-qPCR	6.35	Spike	P	0 - 25

Reference Method: SOP-PCR-1.3
Batch ID: P353626

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2030524	GFD-purified-qPCR	16.0	Spike	P	0 - 25

Reference Method: SOP-PCR-1.4
Batch ID: P352839

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2026987	BacR-qPCR	16.2	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: 2029548
Field Sample ID: 20030793

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	94.6	P	50 - 300
SOP-PCR-1.2	SKETA-qPCR	104	P	50 - 300
SOP-PCR-1.2	SKETA-qPCR	104	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	94.2	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	102							25.5	
	Chlorophyll-a, Uncorrected								25.5	
SOP-PCR-1.0	HF183-qPCR	130	110		99.4	102	96.0			6.35
					79.5					
SOP-PCR-1.2	GULL2-qPCR	179	131	123	84.1	121	84.1			
		102								
SOP-PCR-1.3	GFD-purified-qPCR	13.2	148		0.0	114	107			16.0
					114					
SOP-PCR-1.4	BacR-qPCR	184	102	311	91.7	88.1	108			16.2
		126								

Reference Method Descriptions

Method / DoH Cert #	Description	Associated Samples
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Reference Method Descriptions

Method / DoH Cert #	Description	<u>Associated Samples</u>
SM 10200 H (mod.) / E31780	Phytoplankton chlorophyll-a corrected, uncorrected, and phaeophytin by spectrophotometry	2029546
SM 9223 B Quanti-Tray / E82502	Escherichia coli by Colilert Quanti-Tray method by ALS Environmental Laboratory in Jacksonville	2029547
SOP-PCR-1.0 / E31780	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	2029548
SOP-PCR-1.2 / E31780	Detection and quantification of coastal bird specific Catellicoccus marimammalium Gull2 genetic marker with quantitative polymerase chain reaction	2029548
SOP-PCR-1.3 / E31780	Detection and quantification of bird specific Helicobacter GFD genetic marker with quantitative polymerase chain reaction	2029548
SOP-PCR-1.4 / E31780	Detection and quantification of Ruminant specific Bacteroidetes BacR genetic marker with quantitative polymerase chain reaction	2029548
SOP-PCR-4.0 / E31780	Preparation of Samples by Filtration and held for qPCR Analysis	2029548

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
SM 10200 H (mod.)	09/28/2018	09/28/2018 14:57	Avril Wood	10/16/2018 10:30	Peterson Janvier	2029546
SM 9223 B Quanti-Tray	09/27/2018	09/27/2018 14:33		09/28/2018 14:59	ALS Environmental, Jax	2029547
SOP-PCR-1.0	09/28/2018	09/28/2018 17:00	Sarah Menz	10/22/2018 11:48	Mailin Lopez	2029548
SOP-PCR-1.2	09/28/2018	09/28/2018 17:00	Sarah Menz	10/26/2018 12:44	Mailin Lopez	2029548
SOP-PCR-1.3	09/28/2018	09/28/2018 17:00	Mailin Lopez	11/01/2018 09:39	Sarah Menz	2029548
SOP-PCR-1.4	09/28/2018	09/28/2018 17:00	Sarah Menz	10/25/2018 12:56	Mailin Lopez	2029548
SOP-PCR-4.0	09/28/2018	09/28/2018 17:00	Angela Smith	10/05/2018 13:59	Sarah Menz	2029548