

Biological Analysis Report

WQAP-2017-07-21-04

Florida Department of Environmental Protection
Central Laboratory
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DOH Accreditation E31780

Overflow Analyses Performed By:
Sanders Laboratories
10090 Bavaria Road
Ft. Myers, FL 33913
DOH Accreditation E85457

Event Description: **SMP: Upper Lemon Bay**

Request ID: **RQ-2017-07-17-05**

Customer: **WQAP**

Project ID: **SMP**

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This Report replaces the previous Report Serial Number: 0101417

Report Comment: Re-reported because microbiology was un-authorized to correct incubation times.

Revision certified by: Cheryl A. Swanson, Program Administrator

Date Certified: 03-JUL-2018 17:17



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: WHIDDEN CREEK-BETWEEN TWO MANGROVES

Collection Date/Time: 07/20/2017 14:00

Field ID: G2SD072017-2

Matrix: W-SURF-SLT

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1915514	SM 10200 H (mod.)	Chlorophyll-a, Corrected	4.7		ug/L	P329353	
		Phaeophytin-a	0.85	I	ug/L	P329353	
		Chlorophyll/Phaeophytin Ratio	1.6			P329353	
		Chlorophyll-a, Uncorrected	5.4		ug/L	P329353	

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: LEMON BAY NEAR INDIAN MOUND PARK

Collection Date/Time: 07/20/2017 11:50

Field ID: G3SD072017-1

Matrix: W-SURF-SLT

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1915515	SOP-PCR-1.0	HF183-qPCR**	94	U	GEU/100 mL	P330498	LCS
	SOP-PCR-1.2	GULL2-qPCR**	1.1E+04	UJ	TSC/100 mL	P330498	MS
	SOP-PCR-1.3	GFD-purified-qPCR**	110	U	TSC/100 mL	P330552	LCS, MS, RPD
	SOP-PCR-1.4	BacR-qPCR**	2.0E+03	U	TSC/100 mL	P330498	LCS
	SOP-PCR-4.0	HF183 - Human Source Marker**	Above Threshold				
1915517	Enterolert/QT	ENTEROCOCCI-ENTEROLERT	1990		MPN/100 mL		
	SM 9222 D	Fecal Coliforms-Membrane Filter	2	U	cfu/100 mL		

Ref. Method and Comment:

SOP-PCR-1.0: No amplification of HF183 marker.

SOP-PCR-1.2: No amplification of GULL2 marker. Quality control failure(s) observed. See QA report for details. Precision data unavailable due to inhibition in QC sample.

SOP-PCR-1.3: No amplification of GFD marker.

SOP-PCR-1.4: No amplification of BACR marker. Precision data unavailable.

Sample Location: SARABY0045FTM

Collection Date/Time: 07/20/2017 10:00

Field ID: SARABY0045FTM

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1920859	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray**	184		MPN/100 mL		

Sample Location: G3SD0064

Collection Date/Time: 07/20/2017 10:45

Field ID: G3SD0064

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1920860	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray**	74		MPN/100 mL		

Quality Assurance Report Method Blank Results

Reference Method: SM 10200 H (mod.)

Batch ID: P329353

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: SOP-PCR-1.0
Batch ID: P330498

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

Reference Method: SOP-PCR-1.2
Batch ID: P330498

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

Reference Method: SOP-PCR-1.3
Batch ID: P330552

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

Reference Method: SOP-PCR-1.4
Batch ID: P330498

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: P329353

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	107		P	84 - 116

Reference Method: SOP-PCR-1.0
Batch ID: P330498

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	93.0	2.05	P/F	50 - 175

Reference Method: SOP-PCR-1.2
Batch ID: P330498

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
GULL2-qPCR	82.2	163	P/P	50 - 300

Reference Method: SOP-PCR-1.3
Batch ID: P330552

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
GFD-purified-qPCR	120	12.9	P/F	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P330498

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

Reference Method: SOP-PCR-1.4
Batch ID: P330498

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
BacR-qPCR	134	31.1	P/F	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
Batch ID: P330498

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
1914272	HF183-qPCR	79.4		P	50 - 175
1914985	HF183-qPCR	78.6		P	50 - 175
1915515	HF183-qPCR	70.8		P	50 - 175
1915994	HF183-qPCR	71.6		P	50 - 175
1915995	HF183-qPCR	69.3		P	50 - 175
1915996	HF183-qPCR	82.1		P	50 - 175
1916673	HF183-qPCR	88.5		P	50 - 175
1917382	HF183-qPCR	84.6		P	50 - 175
1922728	HF183-qPCR	81.7		P	50 - 175
1922729	HF183-qPCR	62.3		P	50 - 175
1922730	HF183-qPCR	90.6		P	50 - 175
1922731	HF183-qPCR	71.8		P	50 - 175
1922804	HF183-qPCR	73.0	67.8	P/P	50 - 175

Reference Method: SOP-PCR-1.2
Batch ID: P330498

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
1915515	GULL2-qPCR	35.9		F	50 - 300
1916673	GULL2-qPCR	45.5		F	50 - 300

Reference Method: SOP-PCR-1.3
Batch ID: P330552

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
1914093	GFD-purified-qPCR	112		P	50 - 175
1915515	GFD-purified-qPCR	62.4		P	50 - 175
1915693	GFD-purified-qPCR	120		P	50 - 175
1916673	GFD-purified-qPCR	135		P	50 - 175
1922635	GFD-purified-qPCR	0.0		F	50 - 175
1923592	GFD-purified-qPCR	221		F	50 - 175
1923820	GFD-purified-qPCR	25.5		F	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P330498

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
1915515	BacR-qPCR	132		P	50 - 175
1916673	BacR-qPCR	116		P	50 - 175

Quality Assurance Report Precision

Quality Assurance Report Precision

Reference Method: SM 10200 H (mod.)
Batch ID: P329353

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
1915347	Chlorophyll-a, Corrected	2.74	Sample	P	0 - 25
1915347	Chlorophyll-a, Uncorrected	2.90	Sample	P	0 - 25
1915347	Chlorophyll/Phaeophytin Ratio	0.0	Sample	P	0 - 25

Reference Method: SOP-PCR-1.0
Batch ID: P330498

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
1922804	HF183-qPCR	7.29	Spike	P	0 - 25

Reference Method: SOP-PCR-1.3
Batch ID: P330552

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
1922635	GFD-purified-qPCR	10000	Spike	F	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: 1915515
Field Sample ID: G3SD072017-1

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	78.8	P	50 - 300
SOP-PCR-1.2	SKETA-qPCR	70.6	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	93.1	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	107						2.74	
	Chlorophyll-a, Uncorrected							2.90	
	Chlorophyll/Phaeophytin Ratio							0.0	
SOP-PCR-1.0	HF183-qPCR	2.05	93.0	84.6	81.7	62.3			7.29
				90.6					
SOP-PCR-1.2	GULL2-qPCR	163	82.2	35.9	45.5				
SOP-PCR-1.3	GFD-purified-qPCR	12.9	120	120	0.0	221			10000
				25.5					
SOP-PCR-1.4	BacR-qPCR	31.1	134	132	116				

Reference Method Descriptions

Method / DoH Cert #	Description	Associated Samples
Enterolert/QT / E85457	Enterococci by MPN method by Sanders Laboratories Overflow Laboratory for SD	1915517
SM 10200 H (mod.) / E31780	Phytoplankton chlorophyll-a corrected, uncorrected, and phaeophytin by spectrophotometry	1915514
SM 9222 D / E85457	Fecal coliforms by membrane filter method by Sanders Laboratories Overflow Laboratory for SD	1915517
SM 9223 Quanti-Tray / E85457	Escherichia coli by Colilert 18 Quanti-Tray method by Sanders Laboratories for South District	1920859, 1920860
SOP-PCR-1.0 / E31780	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	1915515
SOP-PCR-1.2 / E31780	Detection and quantification of coastal bird specific Catellicoccus marimammalium Gull2 genetic marker with quantitative polymerase chain reaction	1915515
SOP-PCR-1.3 / E31780	Detection and quantification of bird specific Helicobacter GFD genetic marker with quantitative polymerase chain reaction	1915515
SOP-PCR-1.4 / E31780	Detection and quantification of Ruminant specific Bacteroidetes BacR genetic marker with quantitative polymerase chain reaction	1915515
SOP-PCR-4.0 / E31780	Preparation of Samples by Filtration and held for qPCR Analysis	1915515

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
Enterolert/QT	07/20/2017	07/20/2017 16:10		07/21/2017 16:30	Sanders-Fort Myers	1915517
SM 10200 H (mod.)	07/21/2017	07/21/2017 12:45	Morgan Lane	07/28/2017 07:20	Kim M. Spencer	1915514
SM 9222 D	07/20/2017	07/20/2017 14:50		07/21/2017 14:30	Sanders-Fort Myers	1915517
SM 9223 Quanti-Tray	07/20/2017	07/20/2017 16:00		07/21/2017 11:00	Sanders-Fort Myers	1920859, 1920860
SOP-PCR-1.0	07/21/2017	07/21/2017 12:00	Puja Jasrotia	08/24/2017 12:20	Puja Jasrotia	1915515
SOP-PCR-1.2	07/21/2017	07/21/2017 12:00	Puja Jasrotia	08/28/2017 09:39	Sarah Menz	1915515
SOP-PCR-1.3	07/21/2017	07/21/2017 12:00	Avril Wood	08/29/2017 11:49	Sarah Menz	1915515
SOP-PCR-1.4	07/21/2017	07/21/2017 12:00	Puja Jasrotia	08/25/2017 09:57	Sarah Menz	1915515
SOP-PCR-4.0	07/21/2017	07/21/2017 12:00	Avril Wood	08/21/2017 11:05	Puja Jasrotia	1915515