

Biological Analysis Report

SE-DIST-2021-05-21-02

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

Event Description: **SMP - 3205B pickup**
Request ID: **RQ-2021-05-10-56**
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
Attn: Drew Liddick

For additional information please contact
Cheryl Swanson - Administrator
Jacqueline Savage - Bench Biology & Aquatic
Toxicity
Puja Jasrotia, Ph.D. - Molecular Biology & Taxonomy
Thekkekalathil Chandrasekhar, Ph.D, QA Officer
Phone (850) 245-8177

Certified by: Cheryl A. Swanson, Program Administrator

Date Certified: 18-JUN-2021 10:58



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, Microbiology and Molecular.

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: Taylor Creek @ 70

Collection Date/Time: 05/20/2021 09:45

Field ID: G1SE0011

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2249271	SM 10200 H (mod.)	Chlorophyll-a, Corrected	210		ug/L	P399250	
		Phaeophytin-a	40		ug/L	P399250	
		Chlorophyll-a, Uncorrected	240		ug/L	P399250	
2249272	SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray	517.2	Q	MPN/100 mL	P398758	
2249273	SOP-PCR-1.0	HF183-qPCR**	3.3E+03	UJ	TSC/100 mL	P397905	SUR
	SOP-PCR-1.2	GULL2-qPCR**	3.6E+03	UJ	TSC/100 mL	P397905	SUR
	SOP-PCR-1.3	GFD-qPCR**	720	UJ	TSC/100 mL	P397905	SUR
	SOP-PCR-1.4	BacR-qPCR**	3.0E+03	U	TSC/100 mL	P397905	
	SOP-PCR-1.5	DG3-qPCR**	2.6E+03	UJ	TSC/100 mL	P397905	SUR

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: Refer to the Lab Analysis Report for an explanation of QC Codes.

SOP-PCR-1.2: Refer to the Lab Analysis Report for an explanation of QC Codes.

SOP-PCR-1.3: Refer to the Lab Analysis Report for an explanation of QC Codes.

SOP-PCR-1.5: Refer to the Lab Analysis Report for an explanation of QC Codes.

Quality Assurance Report Method Blank Results

Reference Method: SM 10200 H (mod.)

Batch ID: P399250

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: SM 9223 Quanti-Tray

Batch ID: P398758

Component	Result	Code	Units
Escherichia Coli-Quanti-Tray	1.0	U	MPN/100 mL

Reference Method: SOP-PCR-1.0

Batch ID: P397905

Component	Result	Code	Units
HF183-qPCR	3.3E+03	U	TSC/100 mL

Reference Method: SOP-PCR-1.2

Batch ID: P397905

Component	Result	Code	Units
GULL2-qPCR	3.6E+03	U	TSC/100 mL

Reference Method: SOP-PCR-1.3

Batch ID: P397905

Component	Result	Code	Units
GFD-qPCR	720	U	TSC/100 mL

Quality Assurance Report Method Blank Results

Reference Method: SOP-PCR-1.4
Batch ID: P397905

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

Reference Method: SOP-PCR-1.5
Batch ID: P397905

Component	Result	Code	Units
DG3-qPCR	2.6E+03	U	TSC/100 mL

Quality Assurance Report Laboratory Control Sample Accuracy

Reference Method: SM 10200 H (mod.)
Batch ID: P399250

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

Reference Method: SOP-PCR-1.0
Batch ID: P397905

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	84.4	108	P/P	50 - 175

Reference Method: SOP-PCR-1.2
Batch ID: P397905

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
GULL2-qPCR	113	99.7	P/P	50 - 300

Reference Method: SOP-PCR-1.3
Batch ID: P397905

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
GFD-qPCR	127	100	P/P	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P397905

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
BacR-qPCR	83.2	77.6	P/P	50 - 175

Reference Method: SOP-PCR-1.5
Batch ID: P397905

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
DG3-qPCR	80.6	76.3	P/P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0

Batch ID: P397905

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2246289	HF183-qPCR	65.9	82.4	P	50 - 175
2246290	HF183-qPCR	73.7		P	50 - 175
2246568	HF183-qPCR	85.4		P/P	50 - 175
2247950	HF183-qPCR	95.6		P	50 - 175
2247951	HF183-qPCR	73.5	82.4	P	50 - 175
2248515	HF183-qPCR	67.7		P	50 - 175
2248516	HF183-qPCR	66.5		P	50 - 175
2248895	HF183-qPCR	65.6		P	50 - 175
2249007	HF183-qPCR	83.9		P	50 - 175
2249008	HF183-qPCR	78.6		P	50 - 175
2249084	HF183-qPCR	86.7		P	50 - 175
2249175	HF183-qPCR	87.6		P	50 - 175
2249273	HF183-qPCR	75.2		P	50 - 175

Reference Method: SOP-PCR-1.2

Batch ID: P397905

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2246568	GULL2-qPCR	50.9	52.3	P/P	50 - 300
2247950	GULL2-qPCR	82.5		P	50 - 300
2247951	GULL2-qPCR	71.9		P	50 - 300
2248516	GULL2-qPCR	93.3		P	50 - 300
2248895	GULL2-qPCR	93.4		P	50 - 300
2249007	GULL2-qPCR	75.5		P	50 - 300
2249008	GULL2-qPCR	75.2		P	50 - 300
2249084	GULL2-qPCR	72.0		P	50 - 300
2249175	GULL2-qPCR	91.8		P	50 - 300
2249273	GULL2-qPCR	67.7		P	50 - 300

Reference Method: SOP-PCR-1.3

Batch ID: P397905

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2246568	GFD-qPCR	92.4	93.6	P/P	50 - 175
2247950	GFD-qPCR	96.4		P	50 - 175
2247951	GFD-qPCR	95.0		P	50 - 175
2248516	GFD-qPCR	94.6		P	50 - 175
2248895	GFD-qPCR	102		P	50 - 175
2249007	GFD-qPCR	89.5		P	50 - 175
2249008	GFD-qPCR	87.6		P	50 - 175
2249084	GFD-qPCR	93.2		P	50 - 175
2249175	GFD-qPCR	101		P	50 - 175
2249273	GFD-qPCR	86.4		P	50 - 175

Reference Method: SOP-PCR-1.4

Batch ID: P397905

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2246289	BacR-qPCR	82.8	64.8	P	50 - 175
2246290	BacR-qPCR	82.2		P	50 - 175
2246568	BacR-qPCR	69.4		P/P	50 - 175
2247950	BacR-qPCR	95.8		P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.4

Batch ID: P397905

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2247951	BacR-qPCR	60.9		P	50 - 175
2248515	BacR-qPCR	59.4		P	50 - 175
2248516	BacR-qPCR	73.5		P	50 - 175
2248895	BacR-qPCR	68.2		P	50 - 175
2249007	BacR-qPCR	84.8		P	50 - 175
2249008	BacR-qPCR	79.8		P	50 - 175
2249084	BacR-qPCR	73.8		P	50 - 175
2249175	BacR-qPCR	83.5		P	50 - 175
2249273	BacR-qPCR	53.6		P	50 - 175

Reference Method: SOP-PCR-1.5

Batch ID: P397905

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2246289	DG3-qPCR	76.7		P	50 - 175
2246290	DG3-qPCR	75.3		P	50 - 175
2246568	DG3-qPCR	65.2	66.5	P/P	50 - 175
2247950	DG3-qPCR	89.6		P	50 - 175
2247951	DG3-qPCR	74.1		P	50 - 175
2248515	DG3-qPCR	72.9		P	50 - 175
2248516	DG3-qPCR	69.3		P	50 - 175
2248895	DG3-qPCR	65.9		P	50 - 175
2249007	DG3-qPCR	75.0		P	50 - 175
2249008	DG3-qPCR	73.1		P	50 - 175
2249084	DG3-qPCR	75.1		P	50 - 175
2249175	DG3-qPCR	77.2		P	50 - 175
2249273	DG3-qPCR	70.3		P	50 - 175

Quality Assurance Report Precision

Reference Method: SM 10200 H (mod.)

Batch ID: P399250

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2249074	Chlorophyll-a, Corrected	1.71	Sample	P	0 - 25
2249074	Chlorophyll-a, Uncorrected	0.351	Sample	P	0 - 25

Reference Method: SM 9223 Quanti-Tray

Batch ID: P398758

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2249103	Escherichia Coli-Quanti-Tray	2.02	Sample	P	0 - 80
2249329	Escherichia Coli-Quanti-Tray	66.0	Sample	P	0 - 80

Quality Assurance Report Precision

Reference Method: SOP-PCR-1.0
Batch ID: P397905

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2246568	HF183-qPCR	3.60	Spike	P	0 - 25

Reference Method: SOP-PCR-1.2
Batch ID: P397905

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2246568	GULL2-qPCR	2.76	Spike	P	0 - 25

Reference Method: SOP-PCR-1.3
Batch ID: P397905

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2246568	GFD-qPCR	1.30	Spike	P	0 - 25

Reference Method: SOP-PCR-1.4
Batch ID: P397905

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2246568	BacR-qPCR	6.85	Spike	P	0 - 25

Reference Method: SOP-PCR-1.5
Batch ID: P397905

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2246568	DG3-qPCR	1.93	Spike	P	0 - 25

* Sample, spike and/or laboratory control sample precision (LCS) is reported.

Quality Assurance Report Surrogates

Lab Sample ID: 2249273
Field Sample ID: G1SE0011

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	47.0	F	50 - 300
SOP-PCR-1.2	SKETA-qPCR	47.2	F	50 - 300
SOP-PCR-1.3	SKETA-qPCR	45.5	F	50 - 300
SOP-PCR-1.4	SKETA-qPCR	51.7	P	50 - 300
SOP-PCR-1.5	SKETA-qPCR	43.5	F	50 - 300

Quality Assurance Report Summary

Ref. Method	Analyte	LCS % Recovery		MS % Recovery			LCS	Precision		MS
								SMP		
SM 10200 H (mod.)	Chlorophyll-a, Corrected	102						1.71		
	Chlorophyll-a, Uncorrected							0.351		
SM 9223 Quanti-Tray	Escherichia Coli-Quanti-Tray							2.02	66.0	
SOP-PCR-1.0	HF183-qPCR	108	84.4	65.9	73.7	85.4				3.60
				82.4						
SOP-PCR-1.2	GULL2-qPCR	99.7	113	50.9	52.3	82.5				2.76
				71.9						
SOP-PCR-1.3	GFD-qPCR	100	127	92.4	93.6	96.4				1.30
				95.0						
SOP-PCR-1.4	BacR-qPCR	77.6	83.2	79.8	73.8	83.5				6.85
				53.6						
SOP-PCR-1.5	DG3-qPCR	76.3	80.6	76.7	75.3	65.2				1.93
				66.5						

Reference Method Descriptions

Method	Description	Associated Samples
SM 10200 H (mod.)	Phytoplankton chlorophyll-a corrected, uncorrected, and phaeophytin by spectrophotometry	2249271
SM 9223 Quanti-Tray	Escherichia coli by Colilert 18 Quanti-Tray method - non-chlorinated water systems	2249272
SOP-PCR-1.0	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	2249273
SOP-PCR-1.2	Detection and quantification of coastal bird specific Catellicoccus marimammalium Gull2 genetic marker with quantitative polymerase chain reaction	2249273
SOP-PCR-1.3	Detection and quantification of bird specific Helicobacter GFD genetic marker with quantitative polymerase chain reaction	2249273
SOP-PCR-1.4	Detection and quantification of Ruminant specific Bacteroidetes BacR genetic marker with quantitative polymerase chain reaction	2249273
SOP-PCR-1.5	Detection and quantification of canine-specific DG3 Bacteroides genetic marker with quantitative polymerase chain reaction	2249273

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
SM 10200 H (mod.)	05/21/2021	05/21/2021 14:58	Carina A. Tull	06/01/2021 15:39	Jessica Fetgatter	2249271
SM 9223 Quanti-Tray	05/21/2021	05/21/2021 16:45	Abigail L Baker	05/22/2021 12:29	Mia Jane Harrell	2249272
SOP-PCR-1.0	05/21/2021	05/21/2021 11:50	DeAsia Armster	05/28/2021 10:39	DeAsia Armster	2249273
SOP-PCR-1.2	05/21/2021	05/21/2021 11:50	DeAsia Armster	06/03/2021 11:14	Mailin Lopez	2249273
SOP-PCR-1.3	05/21/2021	05/21/2021 11:50	DeAsia Armster	06/03/2021 13:39	Mailin Lopez	2249273
SOP-PCR-1.4	05/21/2021	05/21/2021 11:50	DeAsia Armster	06/01/2021 14:56	DeAsia Armster	2249273
SOP-PCR-1.5	05/21/2021	05/21/2021 11:50	DeAsia Armster	06/02/2021 12:25	DeAsia Armster	2249273