

Biological Analysis Report

NE-DIST-2020-09-04-01

Florida Department of Environmental Protection
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DOH Accreditation E31780

Overflow Analyses Performed By:
ALS Environmental - Jacksonville
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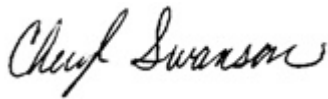
Event Description: **LSJR West**
Request ID: **RQ-2020-08-31-36**
Customer: **NE-DIST**
Project ID: **SMP**

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Certified by: Cheryl A. Swanson, Program Administrator

Date Certified: 02-OCT-2020 12:00



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: Trout River at Garden St

Collection Date/Time: 09/03/2020 09:15

Field ID: 20030751

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2194500	SM 10200 H (mod.)	Chlorophyll-a, Corrected	1.6	U	ug/L	P387266	
		Phaeophytin-a	1.8	U	ug/L	P387266	
		Chlorophyll-a, Uncorrected	1.2	U	ug/L	P387266	
2194508	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	411		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.

Sample Location: Mcgirts Creek at Old Plank Road

Collection Date/Time: 09/03/2020 09:55

Field ID: 20030346

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2194499	SM 10200 H (mod.)	Chlorophyll-a, Corrected	1.6	U	ug/L	P387266	
		Phaeophytin-a	1.8	U	ug/L	P387266	
		Chlorophyll-a, Uncorrected	1.2	U	ug/L	P387266	
2194507	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	649		MPN/100 mL		
2194514	SOP-PCR-1.0	HF183-purified-qPCR**	52	I J	GEU/100 mL	P387144	LCS, SUR
	SOP-PCR-1.4	BacR-purified-qPCR**	3.52E+04	J	TSC/100 mL	P387144	LCS, SUR
	SOP-PCR-1.5	DG3-purified-qPCR**	520	I J	TSC/100 mL	P387144	LCS, 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: Precision data unavailable for Analysis Batch A101301. Refer to the Lab Analysis Report for an explanation of QC Codes.

SOP-PCR-1.4: Precision data unavailable for Analysis Batch A101302. Refer to the Lab Analysis Report for an explanation of QC Codes.

SOP-PCR-1.5: Precision data unavailable for Analysis Batch A101303. Refer to the Lab Analysis Report for an explanation of QC Codes.

Sample Location: Wills Br at Watergate Rd.

Collection Date/Time: 09/03/2020 11:55

Field ID: G2NE1183

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2194502	SM 10200 H (mod.)	Chlorophyll-a, Corrected	2.4	I	ug/L	P387266	
		Phaeophytin-a	0.90	U	ug/L	P387266	
		Chlorophyll-a, Uncorrected	2.7		ug/L	P387266	
2194509	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	206		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.

Sample Location: Yellow Water Cr at Sr 228

Collection Date/Time: 09/03/2020 11:15

Field ID: 20030549

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2194501	SM 10200 H (mod.)	Chlorophyll-a, Corrected	2.8	U	ug/L	P387266	
		Phaeophytin-a	3.0	U	ug/L	P387266	
		Chlorophyll-a, Uncorrected	2.0	U	ug/L	P387266	

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.

Quality Assurance Report Method Blank Results

Reference Method: SM 10200 H (mod.)
Batch ID: P387266

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: P387144

Component	Result	Code	Units
HF183-purified-qPCR	47	U	GEU/100 mL

Reference Method: SOP-PCR-1.4
Batch ID: P387144

Component	Result	Code	Units
BacR-purified-qPCR	510	U	TSC/100 mL

Reference Method: SOP-PCR-1.5
Batch ID: P387144

Component	Result	Code	Units
DG3-purified-qPCR	430	U	TSC/100 mL

Quality Assurance Report Laboratory Control Sample Accuracy

Reference Method: SM 10200 H (mod.)
Batch ID: P387266

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	92.6		P	84 - 116

Reference Method: SOP-PCR-1.0
Batch ID: P387144

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-purified-qPCR	107	16.3	P/F	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P387144

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
BacR-purified-qPCR	77.8	9.78	P/F	50 - 175

Reference Method: SOP-PCR-1.5
Batch ID: P387144

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
DG3-purified-qPCR	61.4	15.8	P/F	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0

Batch ID: P387144

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2194514	HF183-purified-qPCR	128		P	50 - 175
2196599	HF183-purified-qPCR	98.9		P	50 - 175
2196600	HF183-purified-qPCR	107		P	50 - 175
2197431	HF183-purified-qPCR	104		P	50 - 175

Reference Method: SOP-PCR-1.4

Batch ID: P387144

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2194514	BacR-purified-qPCR	63.9		P	50 - 175
2196600	BacR-purified-qPCR	58.9		P	50 - 175

Reference Method: SOP-PCR-1.5

Batch ID: P387144

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2194514	DG3-purified-qPCR	80.4		P	50 - 175
2196600	DG3-purified-qPCR	69.1		P	50 - 175

Quality Assurance Report Surrogates

Lab Sample ID: 2194514

Field Sample ID: 20030346

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	10.6	F	50 - 300
SOP-PCR-1.4	SKETA-qPCR	11.2	F	30 - 300
SOP-PCR-1.5	SKETA-qPCR	9.73	F	50 - 300

Quality Assurance Report Summary

Ref. Method	Analyte	LCS % Recovery		MS % Recovery			LCS	Precision SMP	MS
SM 10200 H (mod.)	Chlorophyll-a, Corrected	92.6							
SOP-PCR-1.0	HF183-purified-qPCR	16.3	107	98.9	107	104			
				128					
SOP-PCR-1.4	BacR-purified-qPCR	9.78	77.8	63.9	58.9				
SOP-PCR-1.5	DG3-purified-qPCR	15.8	61.4	80.4	69.1				

Reference Method Descriptions

Method	Description	Associated Samples
SM 10200 H (mod.)	Phytoplankton chlorophyll-a corrected, uncorrected, and phaeophytin by spectrophotometry	2194499, 2194500, 2194501, 2194502
SM 9223 B Quanti-Tray	E. coli by Colilert/QT by ALS in Jacksonville	2194507, 2194508, 2194509
SOP-PCR-1.0	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	2194514
SOP-PCR-1.4	Detection and quantification of Ruminant specific Bacteroidetes BacR genetic marker with quantitative polymerase chain reaction	2194514

Reference Method Descriptions

Method	Description	<u>Associated Samples</u>
SOP-PCR-1.5	Detection and quantification of canine-specific DG3 Bacteroides genetic marker with quantitative polymerase chain reaction	2194514

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
SM 10200 H (mod.)	09/04/2020	09/04/2020 14:15	Jessica Sherman	09/11/2020 09:08	Yuliya Danyuk	2194499, 2194500, 2194501, 2194502
SM 9223 B Quanti-Tray	09/03/2020	09/03/2020 16:04		09/04/2020 16:05	ALS Environmental, Jax	2194507, 2194508, 2194509
SOP-PCR-1.0	09/04/2020	09/04/2020 14:00	Lauren Campbell	09/30/2020 13:36	Lauren Campbell	2194514
SOP-PCR-1.4	09/04/2020	09/04/2020 14:00	Lauren Campbell	09/30/2020 14:31	Lauren Campbell	2194514
SOP-PCR-1.5	09/04/2020	09/04/2020 14:00	Lauren Campbell	09/30/2020 15:34	Lauren Campbell	2194514