

Biological Analysis Report

NE-DIST-2019-08-21-02

Florida Department of Environmental Protection
Central Laboratory
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DOH Accreditation E31780

Overflow Analyses Performed By:
ALS Environmental - Jacksonville
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DOH Accreditation E82502

Event Description: **Bmap 2**
Request ID: **RQ-2019-08-19-15**
Customer: **NE-DIST**
Project ID: **SMP**

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Date Certified: 17-SEP-2019 16:16



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 R @ DINSMORE BOAT RAMP DOCK
 NR US 1**

Collection Date/Time: 08/20/2019 09:20

Field ID: 20030123

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2113219	SM 10200 H (mod.)	Chlorophyll-a, Corrected	15		ug/L	P370686	
		Phaeophytin-a	1.8	U	ug/L	P370686	
		Chlorophyll-a, Uncorrected	16		ug/L	P370686	
2113223	ASTM D650399	ENTEROCOCCI-ENTEROLERT	580		MPN/100 mL		
2113228	SOP-PCR-1.0	HF183-qPCR**	280	U	GEU/100 mL	P369163	

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: TROUT RIVER AT OLD KINGS RD

Collection Date/Time: 08/20/2019 10:00

Field ID: 20030753

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2113220	SM 10200 H (mod.)	Chlorophyll-a, Corrected	2.1	U	ug/L	P370686	
		Phaeophytin-a	2.2	U	ug/L	P370686	
		Chlorophyll-a, Uncorrected	1.5	U	ug/L	P370686	
2113224	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	488		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: MONCRIEF CR @ 21ST ST

Collection Date/Time: 08/20/2019 10:25

Field ID: 20030897

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2113226	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	1050		MPN/100 mL		

Sample Location: MONCRIEF CR AT MONCRIEF ROAD

Collection Date/Time: 08/20/2019 10:40

Field ID: 20030726

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2113218	SM 10200 H (mod.)	Chlorophyll-a, Corrected	3.2		ug/L	P369849	
		Phaeophytin-a	2.8		ug/L	P369849	
		Chlorophyll-a, Uncorrected	5.0		ug/L	P369849	
2113222	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	34		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: BIG FISHWEIR CR SF @ HAMILTON ST

Collection Date/Time: 08/20/2019 11:20

Field ID: 20030801

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2113225	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	1410		MPN/100 mL		
2113229	SOP-PCR-1.0	HF183-qPCR**	220	I	GEU/100 mL	P369163	

Sample Location: LITTLE FISHWEIR AT OAK ST.

Collection Date/Time: 08/20/2019 11:45

Field ID: 20030953

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2113217	SM 10200 H (mod.)	Chlorophyll-a, Corrected	4.7		ug/L	P369849	
		Phaeophytin-a	0.90	U	ug/L	P369849	
		Chlorophyll-a, Uncorrected	4.9		ug/L	P369849	
2113221	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	1300		MPN/100 mL		
2113227	SOP-PCR-1.0	HF183-qPCR**	200	I	GEU/100 mL	P369163	

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: P369849

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 10200 H (mod.)

Batch ID: P370686

Component	Result	Code	Units
Chlorophyll-a, Corrected	0.89	U	ug/L
Chlorophyll-a, Uncorrected	0.65	U	ug/L
Phaeophytin-a	0.98	U	ug/L

Reference Method: SOP-PCR-1.0

Batch ID: P369163

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

Quality Assurance Report Laboratory Control Sample Accuracy

Reference Method: SM 10200 H (mod.)

Batch ID: P369849

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	87.6		P	84 - 116

Reference Method: SM 10200 H (mod.)

Batch ID: P370686

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	87.4		P	84 - 116

Reference Method: SOP-PCR-1.0

Batch ID: P369163

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	133	150	P/P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0

Batch ID: P369163

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2112421	HF183-qPCR	119	128	P/P	50 - 175
2112429	HF183-qPCR	109		P	50 - 175
2112732	HF183-qPCR	107		P	50 - 175
2112733	HF183-qPCR	94.1		P	50 - 175
2112870	HF183-qPCR	96.2		P	50 - 175
2113066	HF183-qPCR	109		P	50 - 175
2113067	HF183-qPCR	113		P	50 - 175
2113068	HF183-qPCR	112		P	50 - 175
2113227	HF183-qPCR	106		P	50 - 175
2113228	HF183-qPCR	103		P	50 - 175
2113229	HF183-qPCR	98.0		P	50 - 175
2113559	HF183-qPCR	87.7		P	50 - 175

Quality Assurance Report Precision

Reference Method: SM 10200 H (mod.)

Batch ID: P369849

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2113136	Chlorophyll-a, Corrected	2.06	Sample	P	0 - 25
2113136	Chlorophyll-a, Uncorrected	1.80	Sample	P	0 - 25

Reference Method: SM 10200 H (mod.)

Batch ID: P370686

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2113428	Chlorophyll-a, Corrected	4.64	Sample	P	0 - 25
2113428	Chlorophyll-a, Uncorrected	0.842	Sample	P	0 - 25

Reference Method: SOP-PCR-1.0

Batch ID: P369163

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2112421	HF183-qPCR	7.40	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: 2113227

Field Sample ID: 20030953

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	91.2	P	50 - 300

Quality Assurance Report Surrogates

Lab Sample ID: 2113228
 Field Sample ID: 20030123

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	95.2	P	50 - 300

Lab Sample ID: 2113229
 Field Sample ID: 20030801

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	91.9	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	87.6			2.06	
	Chlorophyll-a, Corrected	87.4			4.64	
	Chlorophyll-a, Uncorrected				1.80	
	Chlorophyll-a, Uncorrected				0.842	
SOP-PCR-1.0	HF183-qPCR	150 133	119 128 109 107			7.40

Reference Method Descriptions

Method / DoH Cert #	Description	Associated Samples
ASTM D650399 / E82502	Enterococci by Enterolert/QT by ALS in Jacksonville	2113223
SM 10200 H (mod.) / E31780	Phytoplankton chlorophyll-a corrected, uncorrected, and phaeophytin by spectrophotometry	2113217, 2113218, 2113219, 2113220
SM 9223 B Quanti-Tray / E82502	E. coli by Colilert/QT by ALS in Jacksonville	2113221, 2113222, 2113224, 2113225, 2113226
SOP-PCR-1.0 / E31780	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	2113227, 2113228, 2113229

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
ASTM D650399	08/20/2019	08/20/2019 13:49		08/21/2019 14:29	ALS Environmental, Jax	2113223
SM 10200 H (mod.)	08/21/2019	08/21/2019 14:01	Yuliya Danyuk	08/28/2019 09:14	Yuliya Danyuk	2113217, 2113218
	08/21/2019	08/21/2019 14:01	Yuliya Danyuk	08/28/2019 09:58	Yuliya Danyuk	2113219, 2113220
SM 9223 B Quanti-Tray	08/20/2019	08/20/2019 14:02		08/21/2019 14:31	ALS Environmental, Jax	2113221, 2113222, 2113224, 2113225, 2113226
SOP-PCR-1.0	08/21/2019	08/21/2019 14:00	Angela Smith	09/12/2019 14:03	Mailin Lopez	2113227, 2113228, 2113229