

Biological Analysis Report

NE-DIST-2018-02-07-01

Florida Department of Environmental Protection
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DOH Accreditation E31780

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

Event Description: **bmap tribs**
Request ID: **RQ-2018-01-29-11**
Customer: **NE-DIST**
Project ID: **SMP**

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Date Certified: 19-JUN-2018 08:07



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: Trout at Old Kings

Collection Date/Time: 02/06/2018 11:05

Field ID: 20030753

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1965302	SM 10200 H (mod.)	Chlorophyll-a, Corrected	0.74	I	ug/L	P340031	
		Phaeophytin-a	0.40	U	ug/L	P340031	
		Chlorophyll-a, Uncorrected	0.90	I	ug/L	P340031	
1965308	SM 9223 B	Escherichia Coli-Quanti-Tray**	355		MPN/100 mL		
1965314	SOP-PCR-4.0	HF183 - Human Source Marker**	Below Threshold				

Sample Location: NW Tributary to Open CR @ Hodges Blvd

Collection Date/Time: 02/06/2018 09:10

Field ID: 20030848

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1965303	SM 10200 H (mod.)	Chlorophyll-a, Corrected	3.2	A	ug/L	P340031	
		Phaeophytin-a	2.4	A	ug/L	P340031	
		Chlorophyll-a, Uncorrected	4.8	A	ug/L	P340031	
1965309	SM 9223 B	Escherichia Coli-Quanti-Tray**	504		MPN/100 mL		
1965315	SOP-PCR-1.0	HF183-qPCR**	94	U	GEU/100 mL	P340391	
	SOP-PCR-1.2	GULL2-qPCR**	5.6E+03	U	TSC/100 mL	P340391	
	SOP-PCR-1.3	GFD-purified-qPCR**	4.5	T	TSC/100 mL	P340348	LCS
	SOP-PCR-1.4	BacR-qPCR**	1.0E+03	U	TSC/100 mL	P340391	
	SOP-PCR-4.0	HF183 - Human Source Marker**	Above Threshold				

Ref. Method and Comment:

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

SOP-PCR-1.2: No amplification of GULL2 marker. Precision data unavailable for Analysis Batch A82390.

SOP-PCR-1.3: Precision data unavailable for Analysis Batch A82407.

SOP-PCR-1.4: No amplification of BACR marker. Precision data unavailable for Analysis Batch A82387.

Sample Location: Goodbys at Plaza Gate Rd.

Collection Date/Time: 02/06/2018 13:40

Field ID: 20030537

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1965304	SM 10200 H (mod.)	Chlorophyll-a, Corrected	1.6	I	ug/L	P340031	
		Phaeophytin-a	0.40	U	ug/L	P340031	
		Chlorophyll-a, Uncorrected	1.6		ug/L	P340031	
1965310	SM 9223 B	Escherichia Coli-Quanti-Tray**	24196	J	MPN/100 mL		
1965316	SOP-PCR-1.0	HF183-qPCR**	140	U	GEU/100 mL	P340391	
	SOP-PCR-1.2	GULL2-qPCR**	8.4E+03	U	TSC/100 mL	P340391	
	SOP-PCR-1.3	GFD-purified-qPCR**	110	T	TSC/100 mL	P340348	LCS
	SOP-PCR-1.4	BacR-qPCR**	1.5E+03	U	TSC/100 mL	P340391	
	SOP-PCR-4.0	HF183 - Human Source Marker**	Above Threshold				

Ref. Method and Comment:

SM 9223 B: All Quanti-Tray wells were positive. Result may be higher than reported.

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

SOP-PCR-1.2: No amplification of GULL2 marker. Precision data unavailable for Analysis Batch A82390.

SOP-PCR-1.3: Precision data unavailable for Analysis Batch A82407.

SOP-PCR-1.4: No amplification of BACR marker. Precision data unavailable for Analysis Batch A82387.

Sample Location: Goodbys Trib above San Clerc Rd.

Collection Date/Time: 02/06/2018 13:05

Field ID: 20030891

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1965305	SM 10200 H (mod.)	Chlorophyll-a, Corrected	3.9	U	ug/L	P340031	
		Phaeophytin-a	0.40		ug/L	P340031	
		Chlorophyll-a, Uncorrected	4.0		ug/L	P340031	
1965311	SM 9223 B	Escherichia Coli-Quanti-Tray**	265		MPN/100 mL		
1965317	SOP-PCR-4.0	HF183 - Human Source Marker**	Below Threshold				

Sample Location: Moncrief Creek at 33rd

Collection Date/Time: 02/06/2018 12:20

Field ID: 20030316

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1965306	SM 10200 H (mod.)	Chlorophyll-a, Corrected	18		ug/L	P340031	
		Phaeophytin-a	2.2		ug/L	P340031	
		Chlorophyll-a, Uncorrected	20		ug/L	P340031	
1965312	SM 9223 B	Escherichia Coli-Quanti-Tray**	428		MPN/100 mL		
1965318	SOP-PCR-1.0	HF183-qPCR**	1.61E+03		GEU/100 mL	P340391	
	SOP-PCR-4.0	HF183 - Human Source Marker**	Above Threshold				

Sample Location: Moncrief at 21st

Collection Date/Time: 02/06/2018 11:45

Field ID: 20030897

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1965307	SM 10200 H (mod.)	Chlorophyll-a, Corrected	0.97	I	ug/L	P340031	
		Phaeophytin-a	0.40	U	ug/L	P340031	
		Chlorophyll-a, Uncorrected	0.98	I	ug/L	P340031	
1965313	SM 9223 B	Escherichia Coli-Quanti-Tray**	880		MPN/100 mL		
1965319	SOP-PCR-1.0	HF183-qPCR**	6.53E+03		GEU/100 mL	P340391	
	SOP-PCR-4.0	HF183 - Human Source Marker**	Above Threshold				

Quality Assurance Report
Method Blank Results

Reference Method: SM 10200 H (mod.)
Batch ID: P340031

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

Reference Method: SOP-PCR-1.0
Batch ID: P340391

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

Reference Method: SOP-PCR-1.2
Batch ID: P340391

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

Quality Assurance Report Method Blank Results

Reference Method: SOP-PCR-1.3
Batch ID: P340348

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

Reference Method: SOP-PCR-1.4
Batch ID: P340391

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: P340031

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	99.2		P	84 - 116

Reference Method: SOP-PCR-1.0
Batch ID: P340391

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	138	113	P/P	50 - 175

Reference Method: SOP-PCR-1.2
Batch ID: P340391

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
GULL2-qPCR	147	65.6	P/P	50 - 300

Reference Method: SOP-PCR-1.3
Batch ID: P340348

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
GFD-purified-qPCR	123	5.85	P/F	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P340391

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
BacR-qPCR	108	155	P/P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
Batch ID: P340391

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
1960726	HF183-qPCR	102		P	50 - 175
1963638	HF183-qPCR	94.3		P	50 - 175
1964775	HF183-qPCR	89.7		P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
Batch ID: P340391

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
1964776	HF183-qPCR	71.2		P	50 - 175
1964919	HF183-qPCR	82.4		P	50 - 175
1965315	HF183-qPCR	86.0		P	50 - 175
1965316	HF183-qPCR	96.5		P	50 - 175
1965318	HF183-qPCR	81.8		P	50 - 175
1965319	HF183-qPCR	93.1		P	50 - 175
1966062	HF183-qPCR	93.4	88.9	P/P	50 - 175
1966298	HF183-qPCR	88.9		P	50 - 175
1967544	HF183-qPCR	112		P	50 - 175
1969521	HF183-qPCR	87.3		P	50 - 175

Reference Method: SOP-PCR-1.2
Batch ID: P340391

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
1960726	GULL2-qPCR	70.5		P	50 - 300
1965315	GULL2-qPCR	50.9		P	50 - 300
1965316	GULL2-qPCR	97.9		P	50 - 300

Reference Method: SOP-PCR-1.3
Batch ID: P340348

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
1960643	GFD-purified-qPCR	133		P	50 - 175
1960726	GFD-purified-qPCR	150		P	50 - 175
1961840	GFD-purified-qPCR	170		P	50 - 175
1965315	GFD-purified-qPCR	91.2		P	50 - 175
1965316	GFD-purified-qPCR	163		P	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P340391

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
1960726	BacR-qPCR	101		P	50 - 175
1965315	BacR-qPCR	100		P	50 - 175
1965316	BacR-qPCR	89.3		P	50 - 175
1967544	BacR-qPCR	87.5		P	50 - 175

Quality Assurance Report Precision

Reference Method: SM 10200 H (mod.)
Batch ID: P340031

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
1965303	Chlorophyll-a, Corrected	3.34	Sample	P	0 - 25
1965303	Chlorophyll-a, Uncorrected	1.92	Sample	P	0 - 25
1965303	Phaeophytin-a	1.21	Sample	P	0 - 25

Quality Assurance Report Precision

Reference Method: SOP-PCR-1.0
 Batch ID: P340391

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
1966062	HF183-qPCR	5.00	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: 1965315
 Field Sample ID: 20030848

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	80.7	P	50 - 300
SOP-PCR-1.2	SKETA-qPCR	92.1	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	88.3	P	50 - 300

Lab Sample ID: 1965316
 Field Sample ID: 20030537

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	99.8	P	50 - 300
SOP-PCR-1.2	SKETA-qPCR	99.2	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	106	P	50 - 300

Lab Sample ID: 1965318
 Field Sample ID: 20030316

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	82.1	P	50 - 300

Lab Sample ID: 1965319
 Field Sample ID: 20030897

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	99.1	P	50 - 300

Quality Assurance Report Summary

Ref. Method	Analyte	LCS % Recovery		MS % Recovery			Precision
				LCS			SMP
SM 10200 H (mod.)	Chlorophyll-a, Corrected	99.2					3.34
	Chlorophyll-a, Uncorrected						1.92
	Phaeophytin-a						1.21
SOP-PCR-1.0	HF183-qPCR	113	138	87.3	102	88.9	5.00
SOP-PCR-1.2	GULL2-qPCR	65.6	147	112	70.5	50.9	97.9
SOP-PCR-1.3	GFD-purified-qPCR	5.85	123	133	150	170	91.2

Quality Assurance Report Summary

Ref. Method	Analyte	LCS % Recovery		MS % Recovery			Precision	
							LCS	SMP
SOP-PCR-1.4	BacR-qPCR	155	108	101	87.5	100		
				89.3				

Reference Method Descriptions

Method / DoH Cert #	Description	Associated Samples
SM 10200 H (mod.) / E31780	Phytoplankton chlorophyll-a corrected, uncorrected, and phaeophytin by spectrophotometry	1965302, 1965303, 1965304, 1965305, 1965306, 1965307
SM 9223 B / E82502	Escherichia coli by Colilert Quanti-Tray method by ALS Environmental Overflow Laboratory for Northeast District	1965308, 1965309, 1965310, 1965311, 1965312, 1965313
SOP-PCR-1.0 / E31780	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	1965315, 1965316, 1965318, 1965319
SOP-PCR-1.2 / E31780	Detection and quantification of coastal bird specific Catellicoccus marimammalium Gull2 genetic marker with quantitative polymerase chain reaction	1965315, 1965316
SOP-PCR-1.3 / E31780	Detection and quantification of bird specific Helicobacter GFD genetic marker with quantitative polymerase chain reaction	1965315, 1965316
SOP-PCR-1.4 / E31780	Detection and quantification of Ruminant specific Bacteroidetes BacR genetic marker with quantitative polymerase chain reaction	1965315, 1965316
SOP-PCR-4.0 / E31780	Preparation of Samples by Filtration and held for qPCR Analysis	1965314, 1965315, 1965316, 1965317, 1965318, 1965319

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
SM 10200 H (mod.)	02/07/2018	02/07/2018 13:20	Cathy Sherrer	02/14/2018 12:40	Jennifer Mihalic	1965302, 1965303, 1965304, 1965305, 1965306, 1965307
SM 9223 B	02/06/2018	02/06/2018 14:33		02/07/2018 15:48	ALS Environmental, Jax	1965308, 1965309, 1965310, 1965311, 1965312, 1965313
SOP-PCR-1.0	02/07/2018	02/07/2018 13:45	Angela Smith	02/28/2018 14:51	Sarah Menz	1965315, 1965316, 1965318, 1965319
SOP-PCR-1.2	02/07/2018	02/07/2018 13:45	Angela Smith	03/01/2018 14:13	Sarah Menz	1965315, 1965316
SOP-PCR-1.3	02/07/2018	02/07/2018 13:45	Sarah Menz	03/02/2018 11:46	Sarah Menz	1965315, 1965316
SOP-PCR-1.4	02/07/2018	02/07/2018 13:45	Angela Smith	03/01/2018 10:40	Sarah Menz	1965315, 1965316
SOP-PCR-4.0	02/07/2018	02/07/2018 13:45	Angela Smith	02/27/2018 12:51	Puja Jasrotia	1965314, 1965315, 1965316, 1965317, 1965318, 1965319