

Biological Analysis Report

NE-DIST-2017-12-19-03

Florida Department of Environmental Protection
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DOH Accreditation E31780

Overflow Analyses Performed By:
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Event Description: **Gville Make up**
Request ID: **RQ-2017-12-04-38**
Customer: **NE-DIST**
Project ID: **SMP**

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Date Certified: 07-FEB-2018 15:43



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: LITTLE HATCHET CR @ SR 24

Collection Date/Time: 12/18/2017 10:00

Field ID: G1NE008312182017

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1954991	SM 10200 H (mod.)	Chlorophyll-a, Corrected	0.81	I	ug/L	P337876	
		Phaeophytin-a	0.40	U	ug/L	P337876	
		Chlorophyll-a, Uncorrected	0.78	I	ug/L	P337876	

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: LITTLE HATCHET CR @ SR 24

Collection Date/Time: 12/18/2017 10:00

Field ID: G1NE008312182017

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1954995	EPA 1603	Escherichia coli-Membrane Filter	240		cfu/100 mL		
1954999	SOP-PCR-1.0	HF183-qPCR**	440	I	GEU/100 mL	P336939	
	SOP-PCR-4.0	HF183 - Human Source Marker**	Below Threshold				

Sample Location: TURKEY CR @ NW 59TH

Collection Date/Time: 12/18/2017 12:05

Field ID: G1NE001812182017

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1954992	SM 10200 H (mod.)	Chlorophyll-a, Corrected	0.55	U	ug/L	P337876	
		Phaeophytin-a	0.65	I	ug/L	P337876	
		Chlorophyll-a, Uncorrected	0.78	I	ug/L	P337876	

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: TURKEY CR @ NW 59TH

Collection Date/Time: 12/18/2017 12:05

Field ID: G1NE001812182017

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1954996	EPA 1603	Escherichia coli-Membrane Filter	470		cfu/100 mL		
1955000	SOP-PCR-1.0	HF183-qPCR**	110	T	GEU/100 mL	P336939	
	SOP-PCR-4.0	HF183 - Human Source Marker**	Above Threshold				

Sample Location: TUMBLIN CR @ SW SW 9TH

Collection Date/Time: 12/18/2017 10:35

Field ID: G1NE089412182017

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1954993	SM 10200 H (mod.)	Chlorophyll-a, Corrected	0.55	U	ug/L	P337876	
		Phaeophytin-a	0.40	U	ug/L	P337876	
		Chlorophyll-a, Uncorrected	0.40	U	ug/L	P337876	

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: TUMBLIN CR @ SW SW 9TH

Collection Date/Time: 12/18/2017 10:35

Field ID: G1NE089412182017

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1954997	EPA 1603	Escherichia coli-Membrane Filter	900	B	cfu/100 mL		
1955001	SOP-PCR-1.0	HF183-qPCR**	10	T	GEU/100 mL	P336939	
	SOP-PCR-4.0	HF183 - Human Source Marker**	Above Threshold				

Sample Location: HAGUE BR TRIB @ US 441 80M W OF 77 TERRACE

Collection Date/Time: 12/18/2017 12:35

Field ID: G1NE006312182017

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1954998	EPA 1603	Escherichia coli-Membrane Filter	45000		cfu/100 mL		
1955002	SOP-PCR-1.0	HF183-purified-qPCR**	13	TJ	GEU/100 mL	P338413	LCS, SUR
	SOP-PCR-4.0	HF183 - Human Source Marker**	Above Threshold				

Ref. Method and Comment:

SOP-PCR-1.0: Quality control failure(s) observed. See QA report for details. Precision data unavailable for Analysis Batch A81668.

Sample Location: HAGUE BR TRIB @ US 441 80M W OF 77 TERRACE

Collection Date/Time: 12/18/2017 12:35

Field ID: G1NE006312182017

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1954994	SM 10200 H (mod.)	Chlorophyll-a, Corrected	240		ug/L	P337876	
		Phaeophytin-a	8.0	U	ug/L	P337876	
		Chlorophyll-a, Uncorrected	200		ug/L	P337876	

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: P337876

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: P336939

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

Reference Method: SOP-PCR-1.0

Batch ID: P338413

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

Quality Assurance Report Laboratory Control Sample Accuracy

Reference Method: SM 10200 H (mod.)
Batch ID: P337876

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	94.6		P	84 - 116

Reference Method: SOP-PCR-1.0
Batch ID: P336939

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	141	104	P/P	50 - 175
HF183-qPCR	164	127	P/P	50 - 175

Reference Method: SOP-PCR-1.0
Batch ID: P338413

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-purified-qPCR	121	10.0	P/F	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
Batch ID: P336939

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
1954866	HF183-qPCR	97.7		P	50 - 175
1954892	HF183-qPCR	96.3		P	50 - 175
1954999	HF183-qPCR	95.2		P	50 - 175
1955000	HF183-qPCR	110	106	P/P	50 - 175
1955001	HF183-qPCR	108		P	50 - 175
1955239	HF183-qPCR	93.9		P	50 - 175
1955240	HF183-qPCR	78.6		P	50 - 175
1955241	HF183-qPCR	90.6		P	50 - 175
1955266	HF183-qPCR	86.9		P	50 - 175
1956283	HF183-qPCR	85.9		P	50 - 175
1956284	HF183-qPCR	80.4		P	50 - 175
1956285	HF183-qPCR	80.8		P	50 - 175

Reference Method: SOP-PCR-1.0
Batch ID: P338413

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
1955002	HF183-purified-qPCR	114		P	50 - 175

Quality Assurance Report Precision

Reference Method: SOP-PCR-1.0
Batch ID: P336939

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
1955000	HF183-qPCR	3.09	Spike	P	0 - 25
1958501	HF183-qPCR	3.82	Spike	P	0 - 25

Quality Assurance Report Precision

* 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: 1954999
 Field Sample ID: G1NE008312182017

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	119	P	50 - 300

Lab Sample ID: 1955000
 Field Sample ID: G1NE001812182017

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	122	P	50 - 300

Lab Sample ID: 1955001
 Field Sample ID: G1NE089412182017

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	122	P	50 - 300

Lab Sample ID: 1955002
 Field Sample ID: G1NE006312182017

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	3.19	F	30 - 300

Quality Assurance Report Summary

Ref. Method	Analyte	LCS % Recovery	MS % Recovery	LCS	Precision SMP	MS
SM 10200 H (mod.)	Chlorophyll-a, Corrected	94.6				
SOP-PCR-1.0	HF183-purified-qPCR	10.0 121	114			
	HF183-qPCR	141 127 104	78.6 97.7 96.3			3.82 3.09
		164	95.2			

Reference Method Descriptions

Method / DoH Cert #	Description	Associated Samples
EPA 1603 / E82001	Escherichia coli by membrane filter method using modified mTEC by Advanced Environmental Gainesville Overflow Laboratories for NED	1954995, 1954996, 1954997, 1954998
SM 10200 H (mod.) / E31780	Phytoplankton chlorophyll-a corrected, uncorrected, and phaeophytin by spectrophotometry	1954991, 1954992, 1954993, 1954994
SOP-PCR-1.0 / E31780	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	1954999, 1955000, 1955001
SOP-PCR-1.0 / E31780	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	1955002
SOP-PCR-4.0 / E31780	Preparation of Samples by Filtration and held for qPCR Analysis	1954999, 1955000, 1955001, 1955002

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
EPA 1603	12/18/2017	12/18/2017 15:48		12/19/2017 17:20	AEL - Gainesville	1954995, 1954996, 1954997, 1954998
SM 10200 H (mod.)	12/19/2017	12/19/2017 13:16	Cathy Sherrer	01/05/2018	Jennifer Mihalic	1954991, 1954992, 1954993, 1954994
SOP-PCR-1.0	12/19/2017	12/19/2017 13:30	Avril Wood	01/09/2018	Sarah Menz	1954999, 1955000, 1955001
	12/19/2017	12/20/2017 09:00	Sarah Menz	01/25/2018	Sarah Menz	1955002
SOP-PCR-4.0	12/19/2017	12/19/2017 13:30	Avril Wood	01/16/2018	Puja Jasrotia	1954999, 1955000, 1955001
	12/19/2017	12/20/2017 09:00	Avril Wood	01/16/2018	Puja Jasrotia	1955002