

Biological Analysis Report

CEN-DIST-2021-10-01-01

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

Event Description: **Tributary to Econlockhatchee SCI**

Request ID: **RQ-2021-09-27-36**

Customer: **CEN-DIST**

Project ID: **SMP**

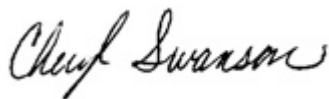
Send Reports to:
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Orlando, FL 32803-3767
Attn: Matt Bearden

For additional information please contact
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Phone (850) 245-8177

Report Comment: The Practical Quantitation Limit associated with the analyte pendimethalin was corrected in the LIMS database. There are no changes from the previous report.

Certified by: Cheryl A. Swanson, Program Administrator

Date Certified: 28-DEC-2021 17:56



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, InvertZool, 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: Econ River Trib @ 170m N of Jay Blanchard Trail

Collection Date/Time: 09/30/2021 13:00

Field ID: G2CE0198

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2279969	SM 10200 H (mod.)	Chlorophyll-a, Corrected	2.9	I	ug/L	P404932	
		Phaeophytin-a	1.8	I	ug/L	P404932	
		Chlorophyll-a, Uncorrected	4.1		ug/L	P404932	
2279970	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	139.6	Q	MPN/100 mL	P404648	RPD
2279972	SOP-PCR-1.0	HF183-qPCR**	2.2E+03	U	TSC/100 mL	P404278	
	SOP-PCR-1.2	GULL2-qPCR**	2.4E+03	UJ	TSC/100 mL	P404278	MS
	SOP-PCR-1.3	GFD-qPCR**	480	U	TSC/100 mL	P404278	
	SOP-PCR-1.4	BacR-qPCR**	2.0E+03	U	TSC/100 mL	P404278	LCS
	SOP-PCR-1.5	DG3-qPCR**	1.7E+03	U	TSC/100 mL	P404278	LCS

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.

SM 9223 B Quanti-Tray: Laboratory duplicate precision is within criteria limits.

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

Sample Location: Econ River Trib @ 170m N of Jay Blanchard Trail

Collection Date/Time: 09/30/2021 13:00

Field ID: G2CE0198

Matrix: BIOLOGICAL

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2279971	SOP-IZ06	Stream Condition Index (SCI)**	41		Points	P406132	
		Macroinvert-FW-Qual-DipnetX20-#Taxa-Rep2**	21		# Taxa	P406132	

Quality Assurance Report Method Blank Results

Reference Method: SM 10200 H (mod.)

Batch ID: P404932

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 B Quanti-Tray

Batch ID: P404648

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

Reference Method: SOP-PCR-1.0

Batch ID: P404278

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

Reference Method: SOP-PCR-1.2

Batch ID: P404278

Component	Result	Code	Units
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Quality Assurance Report Method Blank Results

Reference Method: SOP-PCR-1.2
 Batch ID: P404278

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

Reference Method: SOP-PCR-1.3
 Batch ID: P404278

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

Reference Method: SOP-PCR-1.4
 Batch ID: P404278

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

Reference Method: SOP-PCR-1.5
 Batch ID: P404278

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: P404932

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	106		P	84 - 116

Reference Method: SOP-PCR-1.0
 Batch ID: P404278

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	71.6	113	P/P	50 - 175

Reference Method: SOP-PCR-1.2
 Batch ID: P404278

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
GULL2-qPCR	133	113	P/P	50 - 300

Reference Method: SOP-PCR-1.3
 Batch ID: P404278

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
GFD-qPCR	97.3	99.5	P/P	50 - 175

Reference Method: SOP-PCR-1.4
 Batch ID: P404278

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
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Quality Assurance Report Laboratory Control Sample Accuracy

Reference Method: SOP-PCR-1.4
Batch ID: P404278

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
BacR-qPCR	98.9	19.2	P/F	50 - 175

Reference Method: SOP-PCR-1.5
Batch ID: P404278

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
DG3-qPCR	72.1	19.7	P/F	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
Batch ID: P404278

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2278676	HF183-qPCR	92.9	86.4	P/P	50 - 175
2278854	HF183-qPCR	86.9		P	50 - 175
2279114	HF183-qPCR	94.5		P	50 - 175
2279176	HF183-qPCR	86.2		P	50 - 175
2279587	HF183-qPCR	83.2		P	50 - 175
2279972	HF183-qPCR	76.6		P	50 - 175
2280472	HF183-qPCR	73.1		P	50 - 175
2281065	HF183-qPCR	75.9		P	50 - 175
2281066	HF183-qPCR	77.5		P	50 - 175
2281067	HF183-qPCR	76.4		P	50 - 175
2281068	HF183-qPCR	77.9		P	50 - 175
2281069	HF183-qPCR	79.3		P	50 - 175
2281070	HF183-qPCR	79.3		P	50 - 175

Reference Method: SOP-PCR-1.2
Batch ID: P404278

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2278676	GULL2-qPCR	79.6	77.1	P/P	50 - 300
2278854	GULL2-qPCR	78.8		P	50 - 300
2279114	GULL2-qPCR	72.2		P	50 - 300
2279176	GULL2-qPCR	82.9		P	50 - 300
2279587	GULL2-qPCR	60.1		P	50 - 300
2279972	GULL2-qPCR	47.9		F	50 - 300

Reference Method: SOP-PCR-1.3
Batch ID: P404278

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2278676	GFD-qPCR	106	99.8	P/P	50 - 175
2278854	GFD-qPCR	100		P	50 - 175
2279114	GFD-qPCR	114		P	50 - 175
2279176	GFD-qPCR	103		P	50 - 175
2279587	GFD-qPCR	90.1		P	50 - 175
2279972	GFD-qPCR	91.3		P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.4

Batch ID: P404278

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2278676	BacR-qPCR	77.5	66.4	P/P	50 - 175
2278854	BacR-qPCR	76.6		P	50 - 175
2279114	BacR-qPCR	74.4		P	50 - 175
2279176	BacR-qPCR	72.6		P	50 - 175
2279587	BacR-qPCR	54.4		P	50 - 175
2279972	BacR-qPCR	53.1		P	50 - 175
2280472	BacR-qPCR	58.4		P	50 - 175

Reference Method: SOP-PCR-1.5

Batch ID: P404278

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2278676	DG3-qPCR	86.2	82.0	P/P	50 - 175
2278854	DG3-qPCR	92.3		P	50 - 175
2279114	DG3-qPCR	80.0		P	50 - 175
2279176	DG3-qPCR	88.1		P	50 - 175
2279587	DG3-qPCR	76.1		P	50 - 175
2279972	DG3-qPCR	77.6		P	50 - 175
2280472	DG3-qPCR	77.4		P	50 - 175

Quality Assurance Report Precision

Reference Method: SM 9223 B Quanti-Tray

Batch ID: P404648

Replicated

Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2280070	Escherichia Coli-Quanti-Tray	87.0	Sample	F	0 - 80

Reference Method: SOP-PCR-1.0

Batch ID: P404278

Replicated

Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2278676	HF183-qPCR	7.27	Spike	P	0 - 25

Reference Method: SOP-PCR-1.2

Batch ID: P404278

Replicated

Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2278676	GULL2-qPCR	3.18	Spike	P	0 - 25

Reference Method: SOP-PCR-1.3

Batch ID: P404278

Replicated

Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2278676	GFD-qPCR	5.53	Spike	P	0 - 25

Quality Assurance Report Precision

Reference Method: SOP-PCR-1.4
Batch ID: P404278

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2278676	BacR-qPCR	15.4	Spike	P	0 - 25

Reference Method: SOP-PCR-1.5
Batch ID: P404278

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2278676	DG3-qPCR	5.00	Spike	P	0 - 25

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

Quality Assurance Report Surrogates

Lab Sample ID: 2279972
Field Sample ID: G2CE0198

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	71.2	P	50 - 300
SOP-PCR-1.2	SKETA-qPCR	79.5	P	50 - 300
SOP-PCR-1.3	SKETA-qPCR	94.0	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	83.9	P	50 - 300
SOP-PCR-1.5	SKETA-qPCR	53.9	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	106							
SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray							87.0	
SOP-PCR-1.0	HF183-qPCR	113	71.6	77.5	76.4	77.9			7.27
				79.3					
SOP-PCR-1.2	GULL2-qPCR	113	133	79.6	77.1	78.8			3.18
				72.2					
SOP-PCR-1.3	GFD-qPCR	99.5	97.3	106	99.8	100			5.53
				114					
SOP-PCR-1.4	BacR-qPCR	98.9	19.2	77.5	66.4	76.6			15.4
				74.4					
SOP-PCR-1.5	DG3-qPCR	19.7	72.1	86.2	82.0	92.3			5.00
				80.0					

Reference Method Descriptions

Method	Description	Associated Samples
SM 10200 H (mod.)	Phytoplankton chlorophyll-a corrected, uncorrected, and phaeophytin by spectrophotometry	2279969
SM 9223 B Quanti-Tray	Escherichia coli by Colilert 18 Quanti-Tray method - non-chlorinated water systems	2279970
SOP-IZ06	No. Taxa of FW macroinverts, qual samp, 20 dipnets.	2279971

Reference Method Descriptions

Method	Description	<u>Associated Samples</u>
SOP-PCR-1.0	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	2279972
SOP-PCR-1.2	Detection and quantification of coastal bird specific Catellicoccus marimammalium Gull2 genetic marker with quantitative polymerase chain reaction	2279972
SOP-PCR-1.3	Detection and quantification of bird specific Helicobacter GFD genetic marker with quantitative polymerase chain reaction	2279972
SOP-PCR-1.4	Detection and quantification of Ruminant specific Bacteroidetes BacR genetic marker with quantitative polymerase chain reaction	2279972
SOP-PCR-1.5	Detection and quantification of canine-specific DG3 Bacteroides genetic marker with quantitative polymerase chain reaction	2279972

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
SM 10200 H (mod.)	10/01/2021	10/01/2021 14:06	Jillian G Hanley	10/11/2021 08:33	Jillian G Hanley	2279969
SM 9223 B Quanti-Tray	10/01/2021	10/01/2021 16:11	Amber S. Eells	10/02/2021 12:38	Carina A. Tull	2279970
SOP-IZ06	10/01/2021	10/26/2021 15:45	William T Broome	11/19/2021 09:01	Lisa Homann	2279971
	10/01/2021	10/26/2021 15:45	William T Broome	12/13/2021 09:01	Lisa Homann	2279971
SOP-PCR-1.0	10/01/2021	10/01/2021 13:25	DeAsia Armster	10/08/2021 13:11	DeAsia Armster	2279972
SOP-PCR-1.2	10/01/2021	10/01/2021 13:25	DeAsia Armster	10/13/2021 14:12	DeAsia Armster	2279972
SOP-PCR-1.3	10/01/2021	10/01/2021 13:25	DeAsia Armster	10/15/2021 08:54	Mailin Lopez	2279972
SOP-PCR-1.4	10/01/2021	10/01/2021 13:25	DeAsia Armster	10/12/2021 09:37	DeAsia Armster	2279972
SOP-PCR-1.5	10/01/2021	10/01/2021 13:25	DeAsia Armster	10/13/2021 09:54	DeAsia Armster	2279972