

Biological Analysis Report

SE-DIST-2024-01-30-03

Florida DEP Laboratory
2600 Blair Stone Road
Tallahassee, FL 32399-2400
DOH Accreditation E31780

Event Description: **SMP - Micco**
Request ID: **RQ-2024-01-29-19**
Customer: **SE-DIST**
Project ID: **SMP**

Send Reports to:
FL Dept. of Environmental Protection
FL Dept. of Environmental Protection
2199 South Rock Road
Fort Pierce, FL 34945
Attn: Andrew Luering

For additional information please contact
Cheryl Swanson - Administrator
Sarah Menz, Ph.D. - Bench Biology & Aquatic
Toxicity
Puja Jasrotia, Ph.D. - Molecular Biology & Taxonomy
Thekkekalathil Chandrasekhar, Ph.D, QA Officer
Phone (850) 245-8177

Certified by: Sarah Menz, Environmental Administrator

Date Certified: 13-MAR-2024 12:40

A handwritten signature in black ink, appearing to be 'Sarah Menz', written in a cursive style.

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, 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: Trout Creek near 1st St

Collection Date/Time: 01/29/2024 12:45

Field ID: G5SE0137

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2464216	SM 10150 B	Chlorophyll-a, Corrected	1.2	I	ug/L	P441040	
		Phaeophytin-a	0.90	U	ug/L	P441040	
		Chlorophyll-a, Uncorrected	1.5	I	ug/L	P441040	
2464221	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	206.4	Q	MPN/100 mL	P440975	
2464226	SOP-PCR-1.0	HF183-qPCR**	3.3E+03	U	TSC/100 mL	P440624	
	SOP-PCR-1.2	GULL2-qPCR**	3.6E+03	U	TSC/100 mL	P440624	
	SOP-PCR-1.3	GFD-qPCR**	5.4E+03		TSC/100 mL	P440624	
	SOP-PCR-1.4	BacR-qPCR**	3.0E+03	U	TSC/100 mL	P440624	
	SOP-PCR-1.5	DG3-qPCR**	2.6E+03	U	TSC/100 mL	P440624	

Ref. Method and Comment:

SM 10150 B: 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: Unnamed Canal North of Shell Pit Rd

Collection Date/Time: 01/29/2024 13:10

Field ID: G5SE0138

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2464217	SM 10150 B	Chlorophyll-a, Corrected	1.6	I	ug/L	P441040	
		Phaeophytin-a	1.5	U	ug/L	P441040	
		Chlorophyll-a, Uncorrected	1.1	I	ug/L	P441040	
2464222	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	727.0	Q	MPN/100 mL	P440975	
2464227	SOP-PCR-1.0	HF183-qPCR**	1.7E+03	U	TSC/100 mL	P440624	
	SOP-PCR-1.2	GULL2-qPCR**	1.8E+03	U	TSC/100 mL	P440624	
	SOP-PCR-1.3	GFD-qPCR**	1.6E+03	I	TSC/100 mL	P440624	
	SOP-PCR-1.4	BacR-qPCR**	1.5E+03	U	TSC/100 mL	P440624	
	SOP-PCR-1.5	DG3-qPCR**	1.3E+03	U	TSC/100 mL	P440624	

Ref. Method and Comment:

SM 10150 B: 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: Unnamed Canal North of Senne Highlands

Collection Date/Time: 01/29/2024 13:30

Field ID: G5SE0139

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2464218	SM 10150 B	Chlorophyll-a, Corrected	2.1	I	ug/L	P441040	
		Phaeophytin-a	1.3	U	ug/L	P441040	
		Chlorophyll-a, Uncorrected	2.1	I	ug/L	P441040	
2464223	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	156.5	Q	MPN/100 mL	P440975	
2464228	SOP-PCR-1.0	HF183-qPCR**	1.1E+03	I	TSC/100 mL	P440624	
	SOP-PCR-1.4	BacR-qPCR**	1.0E+03	U	TSC/100 mL	P440624	
	SOP-PCR-1.5	DG3-qPCR**	860	U	TSC/100 mL	P440624	

Ref. Method and Comment:

SM 10150 B: 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: Micco Ditches Weir Off Dottie

Collection Date/Time: 01/29/2024 13:50

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2464215	SM 10150 B	Chlorophyll-a, Corrected	19		ug/L	P441040	
		Phaeophytin-a	6.5		ug/L	P441040	
		Chlorophyll-a, Uncorrected	23		ug/L	P441040	
2464220	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	39.9	Q	MPN/100 mL	P440975	
2464225	SOP-PCR-1.0	HF183-qPCR**	1.7E+03	U	TSC/100 mL	P440624	
	SOP-PCR-1.2	GULL2-qPCR**	1.8E+03	U	TSC/100 mL	P440624	
	SOP-PCR-1.3	GFD-qPCR**	880	I	TSC/100 mL	P440624	
	SOP-PCR-1.4	BacR-qPCR**	1.5E+03	U	TSC/100 mL	P440624	
	SOP-PCR-1.5	DG3-qPCR**	1.3E+03	U	TSC/100 mL	P440624	

SM 10150 B: 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.

Collection Date/Time: 01/29/2024 11:30

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2464219	SM 10150 B	Chlorophyll-a, Corrected	3.4		ug/L	P441040	
		Phaeophytin-a	1.2	I	ug/L	P441040	
		Chlorophyll-a, Uncorrected	4.3		ug/L	P441040	
2464224	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	86.2	Q	MPN/100 mL	P440975	
2464229	SOP-PCR-1.0	HF183-qPCR**	6.6E+03	U	TSC/100 mL	P440624	
	SOP-PCR-1.2	GULL2-qPCR**	7.2E+03	U	TSC/100 mL	P440624	
	SOP-PCR-1.3	GFD-qPCR**	6.8E+03	I	TSC/100 mL	P440624	
	SOP-PCR-1.4	BacR-qPCR**	6.1E+03	U	TSC/100 mL	P440624	
	SOP-PCR-1.5	DG3-qPCR**	5.1E+03	U	TSC/100 mL	P440624	

SM 10150 B: 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

Batch ID: P441040

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

Batch ID: P440975

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

Batch ID: P440624

Component	Result	Code	Units
-----------	--------	------	-------

Quality Assurance Report
Method Blank Results

Reference Method: SOP-PCR-1.0
Batch ID: P440624

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

Reference Method: SOP-PCR-1.2
Batch ID: P440624

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

Reference Method: SOP-PCR-1.3
Batch ID: P440624

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

Reference Method: SOP-PCR-1.4
Batch ID: P440624

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

Reference Method: SOP-PCR-1.5
Batch ID: P440624

Component	Result	Code	Units
DG3-qPCR	2.6E+03	U	TSC/100 mL

Quality Assurance Report
Laboratory Control Sample Accuracy

Reference Method: SM 10150 B
Batch ID: P441040

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	98.1		P	84 - 116

Reference Method: SOP-PCR-1.0
Batch ID: P440624

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	88.8	87.7	P/P	50 - 175

Reference Method: SOP-PCR-1.2
Batch ID: P440624

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
GULL2-qPCR	102	84.5	P/P	50 - 175

Reference Method: SOP-PCR-1.3
Batch ID: P440624

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
-----------	---------	---------	-----------	----------------

Quality Assurance Report
Laboratory Control Sample Accuracy

Reference Method: SOP-PCR-1.3
Batch ID: P440624

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
GFD-qPCR	95.1	86.2	P/P	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P440624

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
BacR-qPCR	90.0	87.7	P/P	50 - 175

Reference Method: SOP-PCR-1.5
Batch ID: P440624

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
DG3-qPCR	101	85.0	P/P	50 - 175

Quality Assurance Report
Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
Batch ID: P440624

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2464107	HF183-qPCR	91.8	91.9	P/P	50 - 175
2464108	HF183-qPCR	109		P	50 - 175
2464193	HF183-qPCR	97.7		P	50 - 175
2464194	HF183-qPCR	92.8		P	50 - 175
2464225	HF183-qPCR	90.6		P	50 - 175
2464226	HF183-qPCR	97.9		P	50 - 175
2464227	HF183-qPCR	86.6		P	50 - 175
2464228	HF183-qPCR	84.1		P	50 - 175
2464229	HF183-qPCR	86.1		P	50 - 175
2464563	HF183-qPCR	94.3		P	50 - 175
2464564	HF183-qPCR	93.1		P	50 - 175
2464565	HF183-qPCR	103		P	50 - 175
2464566	HF183-qPCR	83.8		P	50 - 175

Reference Method: SOP-PCR-1.2
Batch ID: P440624

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2464107	GULL2-qPCR	85.6	91.9	P/P	50 - 175
2464108	GULL2-qPCR	78.4		P	50 - 175
2464193	GULL2-qPCR	66.1		P	50 - 175
2464194	GULL2-qPCR	91.5		P	50 - 175
2464225	GULL2-qPCR	85.2		P	50 - 175
2464226	GULL2-qPCR	86.9		P	50 - 175
2464227	GULL2-qPCR	94.4		P	50 - 175
2464229	GULL2-qPCR	93.3		P	50 - 175
2464564	GULL2-qPCR	89.8		P	50 - 175
2464565	GULL2-qPCR	94.5		P	50 - 175
2464566	GULL2-qPCR	95.0		P	50 - 175

Quality Assurance Report
Matrix Spike Accuracy

Reference Method: SOP-PCR-1.3
Batch ID: P440624

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2464107	GFD-qPCR	106	112	P/P	50 - 175
2464108	GFD-qPCR	106		P	50 - 175
2464193	GFD-qPCR	107		P	50 - 175
2464194	GFD-qPCR	108		P	50 - 175
2464225	GFD-qPCR	93.5		P	50 - 175
2464226	GFD-qPCR	110		P	50 - 175
2464227	GFD-qPCR	108		P	50 - 175
2464229	GFD-qPCR	110		P	50 - 175
2464564	GFD-qPCR	104		P	50 - 175
2464565	GFD-qPCR	107		P	50 - 175
2464566	GFD-qPCR	100		P	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P440624

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2464107	BacR-qPCR	89.0	90.8	P/P	50 - 175
2464108	BacR-qPCR	88.7		P	50 - 175
2464193	BacR-qPCR	76.1		P	50 - 175
2464194	BacR-qPCR	86.5		P	50 - 175
2464225	BacR-qPCR	77.7		P	50 - 175
2464226	BacR-qPCR	87.1		P	50 - 175
2464227	BacR-qPCR	77.1		P	50 - 175
2464228	BacR-qPCR	82.5		P	50 - 175
2464229	BacR-qPCR	82.8		P	50 - 175
2464563	BacR-qPCR	87.6		P	50 - 175
2464564	BacR-qPCR	87.9		P	50 - 175
2464565	BacR-qPCR	84.9		P	50 - 175
2464566	BacR-qPCR	81.2		P	50 - 175

Reference Method: SOP-PCR-1.5
Batch ID: P440624

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2464107	DG3-qPCR	95.4	97.7	P/P	50 - 175
2464108	DG3-qPCR	97.9		P	50 - 175
2464193	DG3-qPCR	82.0		P	50 - 175
2464194	DG3-qPCR	93.7		P	50 - 175
2464225	DG3-qPCR	91.1		P	50 - 175
2464226	DG3-qPCR	99.2		P	50 - 175
2464227	DG3-qPCR	95.9		P	50 - 175
2464228	DG3-qPCR	91.9		P	50 - 175
2464229	DG3-qPCR	94.9		P	50 - 175
2464563	DG3-qPCR	95.4		P	50 - 175
2464564	DG3-qPCR	94.4		P	50 - 175
2464565	DG3-qPCR	83.9		P	50 - 175
2464566	DG3-qPCR	90.8		P	50 - 175

Quality Assurance Report
Precision

Quality Assurance Report Precision

Reference Method: SM 10150 B
Batch ID: P441040

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2464303	Chlorophyll-a, Corrected	1.06	Sample	P	0 - 25
2464303	Chlorophyll-a, Uncorrected	2.07	Sample	P	0 - 25

Reference Method: SOP-PCR-1.0
Batch ID: P440624

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2464107	HF183-qPCR	0.120	Spike	P	0 - 25

Reference Method: SOP-PCR-1.2
Batch ID: P440624

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2464107	GULL2-qPCR	7.05	Spike	P	0 - 25

Reference Method: SOP-PCR-1.3
Batch ID: P440624

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2464107	GFD-qPCR	5.58	Spike	P	0 - 25

Reference Method: SOP-PCR-1.4
Batch ID: P440624

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2464107	BacR-qPCR	2.05	Spike	P	0 - 25

Reference Method: SOP-PCR-1.5
Batch ID: P440624

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2464107	DG3-qPCR	2.33	Spike	P	0 - 25

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

Quality Assurance Report Surrogates

Lab Sample ID: 2464225
Field Sample ID: G5SE0063

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	82.0	P	50 - 300
SOP-PCR-1.2	SKETA-qPCR	118	P	50 - 300

Quality Assurance Report
Surrogates

Lab Sample ID: 2464225
Field Sample ID: G5SE0063

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.3	SKETA-qPCR	106	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	98.2	P	50 - 300
SOP-PCR-1.5	SKETA-qPCR	98.5	P	50 - 300

Lab Sample ID: 2464226
Field Sample ID: G5SE0137

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	86.9	P	50 - 300
SOP-PCR-1.2	SKETA-qPCR	139	P	50 - 300
SOP-PCR-1.3	SKETA-qPCR	121	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	108	P	50 - 300
SOP-PCR-1.5	SKETA-qPCR	106	P	50 - 300

Lab Sample ID: 2464227
Field Sample ID: G5SE0138

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	89.6	P	50 - 300
SOP-PCR-1.2	SKETA-qPCR	126	P	50 - 300
SOP-PCR-1.3	SKETA-qPCR	115	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	107	P	50 - 300
SOP-PCR-1.5	SKETA-qPCR	101	P	50 - 300

Lab Sample ID: 2464228
Field Sample ID: G5SE0139

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	94.7	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	102	P	50 - 300
SOP-PCR-1.5	SKETA-qPCR	109	P	50 - 300

Lab Sample ID: 2464229
Field Sample ID: G5SE0140

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	91.3	P	50 - 300
SOP-PCR-1.2	SKETA-qPCR	131	P	50 - 300
SOP-PCR-1.3	SKETA-qPCR	121	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	106	P	50 - 300
SOP-PCR-1.5	SKETA-qPCR	109	P	50 - 300

Quality Assurance Report
Summary

Ref. Method	Analyte	LCS % Recovery		MS % Recovery			Precision	
							LCS	MS
SM 10150 B	ChlorophyllI-a, Corrected	98.1					1.06	
	ChlorophyllI-a, Uncorrected						2.07	
SOP-PCR-1.0	HF183-qPCR	87.7	88.8	91.8	91.9	109		0.120
				97.7				

Quality Assurance Report

Summary

Ref. Method	Analyte	LCS % Recovery		MS % Recovery			Precision	MS
							SMP	
					LCS			
SOP-PCR-1.2	GULL2-qPCR	84.5	102	85.6	91.9	78.4		7.05
				66.1				
SOP-PCR-1.3	GFD-qPCR	86.2	95.1	93.5	106	112		5.58
				106				
SOP-PCR-1.4	BacR-qPCR	90.0	87.7	77.1	82.5	82.8		2.05
				87.6				
SOP-PCR-1.5	DG3-qPCR	85.0	101	95.4	97.7	97.9		2.33
				82.0				

Reference Method Descriptions

Method	Description	Associated Samples
SM 10150 B	Phytoplankton chlorophyll-a corrected, uncorrected, and phaeophytin by spectrophotometry	2464215, 2464216, 2464217, 2464218, 2464219
SM 9223 B Quanti-Tray	Escherichia coli by Colilert 18 Quanti-Tray method - non-chlorinated water systems	2464220, 2464221, 2464222, 2464223, 2464224
SOP-PCR-1.0	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	2464225, 2464226, 2464227, 2464228, 2464229
SOP-PCR-1.2	Detection and quantification of coastal bird specific Catellicoccus marimammalium Gull2 genetic marker with quantitative polymerase chain reaction	2464225, 2464226, 2464227, 2464229
SOP-PCR-1.3	Detection and quantification of bird specific Helicobacter GFD genetic marker with quantitative polymerase chain reaction	2464225, 2464226, 2464227, 2464229
SOP-PCR-1.4	Detection and quantification of Ruminant specific Bacteroidetes BacR genetic marker with quantitative polymerase chain reaction	2464225, 2464226, 2464227, 2464228, 2464229
SOP-PCR-1.5	Detection and quantification of canine-specific DG3 Bacteroides genetic marker with quantitative polymerase chain reaction	2464225, 2464226, 2464227, 2464228, 2464229

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
SM 10150 B	01/30/2024	01/30/2024 12:58	Keanu Thijm	02/05/2024 10:33	Megan Higbee	2464215, 2464216, 2464217, 2464218, 2464219
SM 9223 B Quanti-Tray	01/30/2024	01/30/2024 16:59	Callie Williams	01/31/2024 13:35	Julia Herten	2464220, 2464221, 2464222, 2464223, 2464224
SOP-PCR-1.0	01/30/2024	01/30/2024 15:06	Lily Gregson	02/20/2024 09:00	Roberta Tatti	2464225
	01/30/2024	01/30/2024 16:20	Lily Gregson	02/20/2024 09:00	Roberta Tatti	2464226, 2464227, 2464228, 2464229
SOP-PCR-1.2	01/30/2024	01/30/2024 15:06	Lily Gregson	02/22/2024 18:06	Madeline Gruver	2464225
	01/30/2024	01/30/2024 16:20	Lily Gregson	02/22/2024 18:06	Madeline Gruver	2464226, 2464227, 2464229
SOP-PCR-1.3	01/30/2024	01/30/2024 15:06	Lily Gregson	02/22/2024 15:21	Madeline Gruver	2464225
	01/30/2024	01/30/2024 16:20	Lily Gregson	02/22/2024 15:21	Madeline Gruver	2464226, 2464227, 2464229
SOP-PCR-1.4	01/30/2024	01/30/2024 15:06	Lily Gregson	02/20/2024 13:27	Roberta Tatti	2464225
	01/30/2024	01/30/2024 16:20	Lily Gregson	02/20/2024 13:27	Roberta Tatti	2464226, 2464227, 2464228, 2464229
SOP-PCR-1.5	01/30/2024	01/30/2024 15:06	Lily Gregson	02/20/2024 14:18	Roberta Tatti	2464225
	01/30/2024	01/30/2024 16:20	Lily Gregson	02/20/2024 14:18	Roberta Tatti	2464226, 2464227, 2464228, 2464229