

Biological Analysis Report

NW-DIST-2021-05-19-01

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

Overflow Analyses Performed By:
University of West Florida
11000 University Parkway
Pensacola, FL 32514
DOH Accreditation E71969

Event Description: **Pace Run**
Request ID: **RQ-2021-05-17-62**
Customer: **NW-DIST**
Project ID: **SMP**

Send Reports to:
FL Dept. of Environmental Protection
160 Government Center
Suite 208
Pensacola, FL 32502-5794
Attn: Cheryl Bunch

For additional information please contact
Cheryl Swanson - Administrator
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Toxicity
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Certified by: Puja Jasrotia, Environmental Administrator

Date Certified: 08-JUN-2021 11:31



NON-CONFORMANCE REPORT INCLUDED

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: Bucket Branch Downstream of Pat Brown Road **Collection Date/Time: 05/18/2021 11:35**

Field ID: G4NW0532

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2248503	SM 10200 H (mod.)	Chlorophyll-a, Corrected	2.8	U	ug/L	P398942	
		Phaeophytin-a	3.0	U	ug/L	P398942	
		Chlorophyll-a, Uncorrected	2.0	U	ug/L	P398942	
2248510	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	145		MPN/100 mL		

Sample Location: Big Coldwater Creek Below CR191

Collection Date/Time: 05/18/2021 12:00

Field ID: G4NW0026

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2248501	SM 10200 H (mod.)	Chlorophyll-a, Corrected	0.82	U	ug/L	P398942	
		Phaeophytin-a	0.90	U	ug/L	P398942	
		Chlorophyll-a, Uncorrected	0.60	U	ug/L	P398942	
2248508	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	41		MPN/100 mL		
2248515	SOP-PCR-1.0	HF183-qPCR**	1.7E+03	U	TSC/100 mL	P397905	
	SOP-PCR-1.4	BacR-qPCR**	2.0E+03	I	TSC/100 mL	P397905	
	SOP-PCR-1.5	DG3-qPCR**	1.3E+03	U	TSC/100 mL	P397905	

Sample Location: Clear Creek Upstream of Pat Brown Road

Collection Date/Time: 05/18/2021 11:15

Field ID: G4NW0360

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2248504	SM 10200 H (mod.)	Chlorophyll-a, Corrected	0.82	U	ug/L	P398678	
		Phaeophytin-a	0.90	U	ug/L	P398678	
		Chlorophyll-a, Uncorrected	0.61	I	ug/L	P398678	
2248511	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	63		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 Coldwater Creek End of Steel Bridge Rd.

Collection Date/Time: 05/18/2021 12:25

Field ID: G4NW0027

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2248502	SM 10200 H (mod.)	Chlorophyll-a, Corrected	0.82	U	ug/L	P398942	
		Phaeophytin-a	0.90	U	ug/L	P398942	
		Chlorophyll-a, Uncorrected	0.60	U	ug/L	P398942	
2248509	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	504		MPN/100 mL		
2248516	SOP-PCR-1.0	HF183-qPCR**	1.7E+03	U	TSC/100 mL	P397905	
	SOP-PCR-1.2	GULL2-qPCR**	1.8E+03	U	TSC/100 mL	P397905	
	SOP-PCR-1.3	GFD-qPCR**	360	U	TSC/100 mL	P397905	
	SOP-PCR-1.4	BacR-qPCR**	1.5E+03	U	TSC/100 mL	P397905	
	SOP-PCR-1.5	DG3-qPCR**	1.3E+03	U	TSC/100 mL	P397905	

Sample Location: Coldwater Creek West Fork Downstream of HWY 87

Collection Date/Time: 05/18/2021 13:15

Field ID: G4NW0017

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2248505	SM 10200 H (mod.)	Chlorophyll-a, Corrected	1.2	U	ug/L	P398678	
		Phaeophytin-a	1.3	U	ug/L	P398678	
		Chlorophyll-a, Uncorrected	0.86	U	ug/L	P398678	
2248512	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	216		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: Coldwater Creek West Fork on CR 178

Collection Date/Time: 05/18/2021 13:35

Field ID: G4NW0016

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2248506	SM 10200 H (mod.)	Chlorophyll-a, Corrected	1.0	UJ	ug/L	P398678	
		Phaeophytin-a	1.1	UJ	ug/L	P398678	
		Chlorophyll-a, Uncorrected	0.75	UJ	ug/L	P398678	
2248513	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	86		MPN/100 mL		

Ref. Method and Comment:

SM 10200 H (mod.): See Non-Conformance Report #8840. 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: Clear Creek Upstream of Hwy 87A

Collection Date/Time: 05/18/2021 14:00

Field ID: G4NW0510

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2248507	SM 10200 H (mod.)	Chlorophyll-a, Corrected	0.82	U	ug/L	P398678	
		Phaeophytin-a	0.90	U	ug/L	P398678	
		Chlorophyll-a, Uncorrected	0.60	U	ug/L	P398678	
2248514	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	10	U	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.

Non-Conformance Report

NCR ID: 8840

Event(s)

NW-DIST-2021-05-19-01

Job(s)

TLH-2021-05-19-70

Sample(s)

2248506

Test(s)

NCR Type: SHIPPING/RECEIVING

NCR Category: Sample Integrity

Observation: Sample container received with loose cap.

Resolution: Data will be qualified as appropriate.

Authorized by/Date: Joshua Ayres 5/26/2021

The Non-Conformance Report details exceptions or problems encountered with the events/jobs/samples/test.

Please address questions to:

Chemistry Colin Wright (850) 245-8085
 Biology Cheryl Swanson (850) 245-8177

Quality Assurance Report

Method Blank Results

Quality Assurance Report Method Blank Results

Reference Method: SM 10200 H (mod.)
Batch ID: P398678

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: P398942

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: SOP-PCR-1.0
Batch ID: P397905

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

Reference Method: SOP-PCR-1.2
Batch ID: P397905

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

Reference Method: SOP-PCR-1.3
Batch ID: P397905

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

Reference Method: SOP-PCR-1.4
Batch ID: P397905

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

Reference Method: SOP-PCR-1.5
Batch ID: P397905

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: P398678

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	104		P	84 - 116

Quality Assurance Report Laboratory Control Sample Accuracy

Reference Method: SM 10200 H (mod.)
Batch ID: P398942

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	101		P	84 - 116

Reference Method: SOP-PCR-1.0
Batch ID: P397905

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	84.4	108	P/P	50 - 175

Reference Method: SOP-PCR-1.2
Batch ID: P397905

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
GULL2-qPCR	113	99.7	P/P	50 - 300

Reference Method: SOP-PCR-1.3
Batch ID: P397905

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
GFD-qPCR	127	100	P/P	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P397905

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
BacR-qPCR	83.2	77.6	P/P	50 - 175

Reference Method: SOP-PCR-1.5
Batch ID: P397905

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
DG3-qPCR	80.6	76.3	P/P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
Batch ID: P397905

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2246289	HF183-qPCR	65.9		P	50 - 175
2246290	HF183-qPCR	73.7		P	50 - 175
2246568	HF183-qPCR	85.4	82.4	P/P	50 - 175
2247950	HF183-qPCR	95.6		P	50 - 175
2247951	HF183-qPCR	73.5		P	50 - 175
2248515	HF183-qPCR	67.7		P	50 - 175
2248516	HF183-qPCR	66.5		P	50 - 175
2248895	HF183-qPCR	65.6		P	50 - 175
2249007	HF183-qPCR	83.9		P	50 - 175
2249008	HF183-qPCR	78.6		P	50 - 175
2249084	HF183-qPCR	86.7		P	50 - 175
2249175	HF183-qPCR	87.6		P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0

Batch ID: P397905

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2249273	HF183-qPCR	75.2		P	50 - 175

Reference Method: SOP-PCR-1.2

Batch ID: P397905

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2246568	GULL2-qPCR	50.9	52.3	P/P	50 - 300
2247950	GULL2-qPCR	82.5		P	50 - 300
2247951	GULL2-qPCR	71.9		P	50 - 300
2248516	GULL2-qPCR	93.3		P	50 - 300
2248895	GULL2-qPCR	93.4		P	50 - 300
2249007	GULL2-qPCR	75.5		P	50 - 300
2249008	GULL2-qPCR	75.2		P	50 - 300
2249084	GULL2-qPCR	72.0		P	50 - 300
2249175	GULL2-qPCR	91.8		P	50 - 300
2249273	GULL2-qPCR	67.7		P	50 - 300

Reference Method: SOP-PCR-1.3

Batch ID: P397905

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2246568	GFD-qPCR	92.4	93.6	P/P	50 - 175
2247950	GFD-qPCR	96.4		P	50 - 175
2247951	GFD-qPCR	95.0		P	50 - 175
2248516	GFD-qPCR	94.6		P	50 - 175
2248895	GFD-qPCR	102		P	50 - 175
2249007	GFD-qPCR	89.5		P	50 - 175
2249008	GFD-qPCR	87.6		P	50 - 175
2249084	GFD-qPCR	93.2		P	50 - 175
2249175	GFD-qPCR	101		P	50 - 175
2249273	GFD-qPCR	86.4		P	50 - 175

Reference Method: SOP-PCR-1.4

Batch ID: P397905

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2246289	BacR-qPCR	82.8		P	50 - 175
2246290	BacR-qPCR	82.2		P	50 - 175
2246568	BacR-qPCR	69.4	64.8	P/P	50 - 175
2247950	BacR-qPCR	95.8		P	50 - 175
2247951	BacR-qPCR	60.9		P	50 - 175
2248515	BacR-qPCR	59.4		P	50 - 175
2248516	BacR-qPCR	73.5		P	50 - 175
2248895	BacR-qPCR	68.2		P	50 - 175
2249007	BacR-qPCR	84.8		P	50 - 175
2249008	BacR-qPCR	79.8		P	50 - 175
2249084	BacR-qPCR	73.8		P	50 - 175
2249175	BacR-qPCR	83.5		P	50 - 175
2249273	BacR-qPCR	53.6		P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.5

Batch ID: P397905

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2246289	DG3-qPCR	76.7		P	50 - 175
2246290	DG3-qPCR	75.3		P	50 - 175
2246568	DG3-qPCR	65.2	66.5	P/P	50 - 175
2247950	DG3-qPCR	89.6		P	50 - 175
2247951	DG3-qPCR	74.1		P	50 - 175
2248515	DG3-qPCR	72.9		P	50 - 175
2248516	DG3-qPCR	69.3		P	50 - 175
2248895	DG3-qPCR	65.9		P	50 - 175
2249007	DG3-qPCR	75.0		P	50 - 175
2249008	DG3-qPCR	73.1		P	50 - 175
2249084	DG3-qPCR	75.1		P	50 - 175
2249175	DG3-qPCR	77.2		P	50 - 175
2249273	DG3-qPCR	70.3		P	50 - 175

Quality Assurance Report Precision

Reference Method: SM 10200 H (mod.)

Batch ID: P398678

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2248842	Chlorophyll-a, Corrected	5.46	Sample	P	0 - 25
2248842	Chlorophyll-a, Uncorrected	4.62	Sample	P	0 - 25

Reference Method: SM 10200 H (mod.)

Batch ID: P398942

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2248327	Chlorophyll-a, Corrected	3.11	Sample	P	0 - 25
2248327	Chlorophyll-a, Uncorrected	3.13	Sample	P	0 - 25
2248327	Phaeophytin-a	3.23	Sample	P	0 - 25

Reference Method: SOP-PCR-1.0

Batch ID: P397905

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2246568	HF183-qPCR	3.60	Spike	P	0 - 25

Reference Method: SOP-PCR-1.2

Batch ID: P397905

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2246568	GULL2-qPCR	2.76	Spike	P	0 - 25

Quality Assurance Report Precision

Reference Method: SOP-PCR-1.3
 Batch ID: P397905

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2246568	GFD-qPCR	1.30	Spike	P	0 - 25

Reference Method: SOP-PCR-1.4
 Batch ID: P397905

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2246568	BacR-qPCR	6.85	Spike	P	0 - 25

Reference Method: SOP-PCR-1.5
 Batch ID: P397905

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2246568	DG3-qPCR	1.93	Spike	P	0 - 25

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

Quality Assurance Report Surrogates

Lab Sample ID: 2248515
 Field Sample ID: G4NW0026

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	95.4	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	91.8	P	50 - 300
SOP-PCR-1.5	SKETA-qPCR	92.3	P	50 - 300

Lab Sample ID: 2248516
 Field Sample ID: G4NW0027

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	93.4	P	50 - 300
SOP-PCR-1.2	SKETA-qPCR	92.8	P	50 - 300
SOP-PCR-1.3	SKETA-qPCR	93.5	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	96.0	P	50 - 300
SOP-PCR-1.5	SKETA-qPCR	90.2	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	104			5.46	
	Chlorophyll-a, Corrected	101			3.11	
	Chlorophyll-a, Uncorrected				4.62	
	Chlorophyll-a, Uncorrected				3.13	
	Phaeophytin-a				3.23	

Quality Assurance Report Summary

Ref. Method	Analyte	LCS % Recovery		MS % Recovery			Precision SMP	MS
SOP-PCR-1.0	HF183-qPCR	108	84.4	65.9	73.7	85.4		3.60
SOP-PCR-1.2	GULL2-qPCR	99.7	113	50.9	52.3	82.5		2.76
SOP-PCR-1.3	GFD-qPCR	100	127	92.4	93.6	96.4		1.30
SOP-PCR-1.4	BacR-qPCR	77.6	83.2	79.8	73.8	83.5		6.85
SOP-PCR-1.5	DG3-qPCR	76.3	80.6	76.7	75.3	65.2		1.93

Reference Method Descriptions

Method	Description	Associated Samples
SM 10200 H (mod.)	Phytoplankton chlorophyll-a corrected, uncorrected, and phaeophytin by spectrophotometry	2248501, 2248502, 2248503, 2248504, 2248505, 2248506, 2248507
SM 9223 B Quanti-Tray	E. coli by Colilert-18/QT by UWF in Pensacola	2248508, 2248509, 2248510, 2248511, 2248512, 2248513, 2248514
SOP-PCR-1.0	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	2248515, 2248516
SOP-PCR-1.2	Detection and quantification of coastal bird specific Catellicoccus marimammalium Gull2 genetic marker with quantitative polymerase chain reaction	2248516
SOP-PCR-1.3	Detection and quantification of bird specific Helicobacter GFD genetic marker with quantitative polymerase chain reaction	2248516
SOP-PCR-1.4	Detection and quantification of Ruminant specific Bacteroidetes BacR genetic marker with quantitative polymerase chain reaction	2248515, 2248516
SOP-PCR-1.5	Detection and quantification of canine-specific DG3 Bacteroides genetic marker with quantitative polymerase chain reaction	2248515, 2248516

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
SM 10200 H (mod.)	05/19/2021	05/19/2021 14:42	Carina A. Tull	05/21/2021 09:58	Jessica Fetgatter	2248501, 2248502, 2248503
	05/19/2021	05/19/2021 14:42	Carina A. Tull	05/24/2021 08:26	Jessica Fetgatter	2248504, 2248505, 2248506, 2248507
SM 9223 B Quanti-Tray	05/18/2021	05/18/2021 16:30		05/19/2021 13:10	UWF Lab	2248508, 2248509, 2248510, 2248511, 2248512, 2248513, 2248514
SOP-PCR-1.0	05/19/2021	05/19/2021 14:10	DeAsia Armster	05/28/2021 10:39	DeAsia Armster	2248515, 2248516
SOP-PCR-1.2	05/19/2021	05/19/2021 14:10	DeAsia Armster	06/03/2021 11:14	Mailin Lopez	2248516
SOP-PCR-1.3	05/19/2021	05/19/2021 14:10	DeAsia Armster	06/03/2021 13:39	Mailin Lopez	2248516
SOP-PCR-1.4	05/19/2021	05/19/2021 14:10	DeAsia Armster	06/01/2021 14:56	DeAsia Armster	2248515, 2248516
SOP-PCR-1.5	05/19/2021	05/19/2021 14:10	DeAsia Armster	06/02/2021 12:25	DeAsia Armster	2248515, 2248516