

Biological Analysis Report

TLH-ROC-2022-08-03-01

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

Event Description: **Run 11**
Request ID: **RQ-2022-07-18-45**
Customer: **TLH-ROC**
Project ID: **SMP**

Send Reports to:
FL Dept. of Environmental Protection
2600 Blair Stone Rd
Tallahassee, FL 32399
Attn: Michael Eckles

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: Cheryl A. Swanson, Program Administrator

Date Certified: 15-SEP-2022 12:01



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: Mule Creek @ Alvie Holmes Rd

Collection Date/Time: 08/03/2022 11:15

Field ID: G3TLHR0017

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2345948	SM 10200 H (mod.)	Chlorophyll-a, Corrected	0.94	I	ug/L	P418131	
		Phaeophytin-a	0.98	U	ug/L	P418131	
		Chlorophyll-a, Uncorrected	0.78	I	ug/L	P418131	
2345959	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	88.4		MPN/100 mL	P417693	
2345962	SOP-PCR-1.0	HF183-qPCR**	1.7E+03	U	TSC/100 mL	P417615	
	SOP-PCR-1.2	GULL2-qPCR**	1.8E+03	U	TSC/100 mL	P417615	
	SOP-PCR-1.3	GFD-qPCR**	360	U	TSC/100 mL	P417615	
	SOP-PCR-1.4	BacR-qPCR**	1.5E+03	U	TSC/100 mL	P417615	
	SOP-PCR-1.5	DG3-qPCR**	1.3E+03	U	TSC/100 mL	P417615	

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: Sandy Creek Upstream Boat Ramp

Collection Date/Time: 08/03/2022 11:40

Field ID: G3TLHR0037

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2345951	Enterolert/QT	Enterococci-Quanti-Tray	41.4	A	MPN/100 mL	P417693	
2345953	SM 9222 D	Fecal Coliforms-Membrane Filter	190	B	cfu/100 mL	P417695	

Ref. Method and Comment:

SM 9222 D: Precision data for QC sample is unavailable due to the colony count falling outside of method acceptable range.

Sample Location: Sandy Creek @ Mouth

Collection Date/Time: 08/03/2022 12:05

Field ID: G3TLHR0051

Matrix: W-SURF-SLT

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2345946	SM 10200 H (mod.)	Chlorophyll-a, Corrected	7.9		ug/L	P418131	
		Phaeophytin-a	2.4	U	ug/L	P418131	
		Chlorophyll-a, Uncorrected	7.8		ug/L	P418131	
2345955	Enterolert/QT	Enterococci-Quanti-Tray	31.0		MPN/100 mL	P417693	
2345957	SM 9222 D	Fecal Coliforms-Membrane Filter	94		cfu/100 mL	P417695	

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 9222 D: Precision data for QC sample is unavailable due to the colony count falling outside of method acceptable range.

Sample Location: Eagle Nest Bayou @ Mouth

Collection Date/Time: 08/03/2022 12:15

Field ID: G3TLHR0041

Matrix: W-SURF-SLT

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2345949	SM 10200 H (mod.)	Chlorophyll-a, Corrected	15	A	ug/L	P418131	
		Phaeophytin-a	2.0	U	ug/L	P418131	
		Chlorophyll-a, Uncorrected	16	A	ug/L	P418131	
2345960	Enterolert/QT	Enterococci-Quanti-Tray	10.0	U	MPN/100 mL	P417693	
2345961	SM 9222 D	Fecal Coliforms-Membrane Filter	22	B	cfu/100 mL	P417695	

Field ID: G3TLHR0041

Matrix: W-SURF-SLT

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
Ref. Method and Comment: SM 10200 H (mod.): Laboratory duplicate result is 2.0 U for Phaeophytin-a. Precision data is unavailable due to the small amount of analyte in the QC sample. SM 9222 D: Laboratory duplicate result is 15 B. Precision data for this duplicate is unavailable due to the colony count falling outside of method acceptable range.							

Sample Location: Eagle Nest Continuous Monitoring

Collection Date/Time: 08/03/2022 12:25

Field ID: G3TLHR0019

Matrix: W-SURF-SLT

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2345952	Enterolert/QT	Enterococci-Quanti-Tray	30.0		MPN/100 mL	P417693	
2345954	SM 9222 D	Fecal Coliforms-Membrane Filter	10	B	cfu/100 mL	P417695	
Ref. Method and Comment: SM 9222 D: Precision data for QC sample is unavailable due to the colony count falling outside of method acceptable range.							

Sample Location: Sandy Creek

Collection Date/Time: 08/03/2022 13:10

Field ID: G3TLHR0001

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2345945	SM 10200 H (mod.)	Chlorophyll-a, Corrected	4.1	U	ug/L	P418131	
		Phaeophytin-a	4.5	U	ug/L	P418131	
		Chlorophyll-a, Uncorrected	3.0	U	ug/L	P418131	
2345950	SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray	156.5		MPN/100 mL	P417693	
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: Enterolert/QT
Batch ID: P417693

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

Reference Method: SM 10200 H (mod.)
Batch ID: P418131

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 9222 D
Batch ID: P417695

Component	Result	Code	Units
Fecal Coliforms-Membrane Filter	2	U	cfu/100 mL

Reference Method: SM 9223 B Quanti-Tray
Batch ID: P417693

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

Quality Assurance Report Method Blank Results

Reference Method: SM 9223 B Quanti-Tray
Batch ID: P417693

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

Reference Method: SOP-PCR-1.0
Batch ID: P417615

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

Reference Method: SOP-PCR-1.2
Batch ID: P417615

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

Reference Method: SOP-PCR-1.3
Batch ID: P417615

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

Reference Method: SOP-PCR-1.4
Batch ID: P417615

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

Reference Method: SOP-PCR-1.5
Batch ID: P417615

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: P418131

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	86.5		P	84 - 116

Reference Method: SOP-PCR-1.0
Batch ID: P417615

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	64.7	83.4	P/P	50 - 175

Reference Method: SOP-PCR-1.2
Batch ID: P417615

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

Quality Assurance Report Laboratory Control Sample Accuracy

Reference Method: SOP-PCR-1.2
Batch ID: P417615

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
GULL2-qPCR	58.8	79.4	P/P	50 - 300

Reference Method: SOP-PCR-1.3
Batch ID: P417615

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
GFD-qPCR	90.4	73.3	P/P	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P417615

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
BacR-qPCR	93.0	83.2	P/P	50 - 175

Reference Method: SOP-PCR-1.5
Batch ID: P417615

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
DG3-qPCR	81.7	95.6	P/P	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
Batch ID: P417615

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2345962	HF183-qPCR	87.9	87.9	P/P	50 - 175
2346162	HF183-qPCR	87.4		P	50 - 175
2346393	HF183-qPCR	82.5		P	50 - 175
2346587	HF183-qPCR	94.6		P	50 - 175
2346598	HF183-qPCR	96.3		P	50 - 175
2346616	HF183-qPCR	102		P	50 - 175
2347375	HF183-qPCR	84.4		P	50 - 175
2347737	HF183-qPCR	92.4		P	50 - 175
2347738	HF183-qPCR	103		P	50 - 175
2347739	HF183-qPCR	101		P	50 - 175
2347740	HF183-qPCR	110		P	50 - 175
2347741	HF183-qPCR	116		P	50 - 175
2347742	HF183-qPCR	101		P	50 - 175

Reference Method: SOP-PCR-1.2
Batch ID: P417615

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2345962	GULL2-qPCR	79.3	78.3	P/P	50 - 300
2346162	GULL2-qPCR	75.8		P	50 - 300
2346393	GULL2-qPCR	90.8		P	50 - 300
2346616	GULL2-qPCR	70.4		P	50 - 300
2347375	GULL2-qPCR	87.6		P	50 - 300
2347737	GULL2-qPCR	62.9		P	50 - 300

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.2

Batch ID: P417615

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2347738	GULL2-qPCR	70.5		P	50 - 300
2347739	GULL2-qPCR	82.3		P	50 - 300
2347740	GULL2-qPCR	88.9		P	50 - 300
2347741	GULL2-qPCR	83.9		P	50 - 300
2347742	GULL2-qPCR	82.5		P	50 - 300

Reference Method: SOP-PCR-1.3

Batch ID: P417615

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2345962	GFD-qPCR	106	102	P/P	50 - 175
2346162	GFD-qPCR	85.7		P	50 - 175
2346393	GFD-qPCR	103		P	50 - 175
2346616	GFD-qPCR	103		P	50 - 175
2347375	GFD-qPCR	93.7		P	50 - 175
2347737	GFD-qPCR	88.4		P	50 - 175
2347738	GFD-qPCR	97.1		P	50 - 175
2347739	GFD-qPCR	92.3		P	50 - 175
2347740	GFD-qPCR	95.1		P	50 - 175
2347741	GFD-qPCR	92.3		P	50 - 175
2347742	GFD-qPCR	93.3		P	50 - 175

Reference Method: SOP-PCR-1.4

Batch ID: P417615

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2345962	BacR-qPCR	71.6	75.4	P/P	50 - 175
2346162	BacR-qPCR	74.2		P	50 - 175
2346393	BacR-qPCR	70.2		P	50 - 175
2346587	BacR-qPCR	93.2		P	50 - 175
2346598	BacR-qPCR	73.8		P	50 - 175
2346616	BacR-qPCR	62.7		P	50 - 175
2347375	BacR-qPCR	86.6		P	50 - 175

Reference Method: SOP-PCR-1.5

Batch ID: P417615

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2345962	DG3-qPCR	94.5	93.2	P/P	50 - 175
2346162	DG3-qPCR	89.6		P	50 - 175
2346393	DG3-qPCR	90.7		P	50 - 175
2346587	DG3-qPCR	102		P	50 - 175
2346598	DG3-qPCR	102		P	50 - 175
2346616	DG3-qPCR	108		P	50 - 175
2347375	DG3-qPCR	102		P	50 - 175
2347737	DG3-qPCR	94.0		P	50 - 175
2347738	DG3-qPCR	98.0		P	50 - 175
2347739	DG3-qPCR	103		P	50 - 175
2347740	DG3-qPCR	103		P	50 - 175
2347741	DG3-qPCR	111		P	50 - 175
2347742	DG3-qPCR	102		P	50 - 175

Quality Assurance Report Precision

Reference Method: Enterolert/QT

Batch ID: P417693

Replicated

Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2345951	Enterococci-Quanti-Tray	47.0	Sample	P	0 - 80

Reference Method: SM 10200 H (mod.)

Batch ID: P418131

Replicated

Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2345949	Chlorophyll-a, Corrected	17.4	Sample	P	0 - 25
2345949	Chlorophyll-a, Uncorrected	19.7	Sample	P	0 - 25

Reference Method: SM 9223 B Quanti-Tray

Batch ID: P417693

Replicated

Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2345356	Escherichia Coli-Quanti-Tray	9.13	Sample	P	0 - 80
2345611	Escherichia Coli-Quanti-Tray	5.58	Sample	P	0 - 80

Reference Method: SOP-PCR-1.0

Batch ID: P417615

Replicated

Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2345962	HF183-qPCR	0.0893	Spike	P	0 - 25

Reference Method: SOP-PCR-1.2

Batch ID: P417615

Replicated

Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2345962	GULL2-qPCR	1.28	Spike	P	0 - 25

Reference Method: SOP-PCR-1.3

Batch ID: P417615

Replicated

Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2345962	GFD-qPCR	4.24	Spike	P	0 - 25

Reference Method: SOP-PCR-1.4

Batch ID: P417615

Replicated

Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2345962	BacR-qPCR	5.23	Spike	P	0 - 25

Reference Method: SOP-PCR-1.5

Batch ID: P417615

Replicated

Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
------------	-----------	-----------	-------------------	-----------	----------------

Quality Assurance Report Precision

Reference Method: SOP-PCR-1.5
 Batch ID: P417615

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2345962	DG3-qPCR	1.35	Spike	P	0 - 25

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

Quality Assurance Report Surrogates

Lab Sample ID: 2345962
 Field Sample ID: G3TLHR0017

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	95.1	P	50 - 300
SOP-PCR-1.0	SKETA-qPCR	101	P	50 - 300
SOP-PCR-1.2	SKETA-qPCR	115	P	50 - 300
SOP-PCR-1.2	SKETA-qPCR	120	P	50 - 300
SOP-PCR-1.3	SKETA-qPCR	120	P	50 - 300
SOP-PCR-1.3	SKETA-qPCR	127	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	115	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	121	P	50 - 300
SOP-PCR-1.5	SKETA-qPCR	99.3	P	50 - 300
SOP-PCR-1.5	SKETA-qPCR	103	P	50 - 300

Quality Assurance Report Summary

Ref. Method	Analyte	LCS % Recovery		MS % Recovery			LCS	Precision		MS
								SMP		
Enterolert/QT	Enterococci-Quanti-Tray							47.0		
SM 10200 H (mod.)	Chlorophyll-a, Corrected	86.5						17.4		
	Chlorophyll-a, Uncorrected							19.7		
SM 9223 B Quanti-Tray	Escherichia Coli-Quanti-Tray							9.13	5.58	
SOP-PCR-1.0	HF183-qPCR	83.4	64.7	87.9	87.9	87.4				0.0893
				82.5						
SOP-PCR-1.2	GULL2-qPCR	79.4	58.8	79.3	78.3	75.8				1.28
				90.8						
SOP-PCR-1.3	GFD-qPCR	73.3	90.4	106	102	85.7				4.24
				103						
SOP-PCR-1.4	BacR-qPCR	83.2	93.0	71.6	75.4	74.2				5.23
				70.2						
SOP-PCR-1.5	DG3-qPCR	95.6	81.7	94.5	93.2	89.6				1.35
				90.7						

Reference Method Descriptions

Method	Description	Associated Samples
Enterolert/QT	Enterococci by Enterolert Quanti-Tray method - non-chlorinated water systems	2345951, 2345952, 2345955, 2345960
SM 10200 H (mod.)	Phytoplankton chlorophyll-a corrected, uncorrected, and phaeophytin by spectrophotometry	2345945, 2345946, 2345948, 2345949
SM 9222 D	Fecal coliforms by membrane filter method - non-chlorinated water systems	2345953, 2345954, 2345957, 2345961

Reference Method Descriptions

Method	Description	<u>Associated Samples</u>
SM 9223 B Quanti-Tray	Escherichia coli by Colilert 18 Quanti-Tray method - non-chlorinated water systems	2345950, 2345959
SOP-PCR-1.0	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	2345962
SOP-PCR-1.2	Detection and quantification of coastal bird specific Catellicoccus marimammalium Gull2 genetic marker with quantitative polymerase chain reaction	2345962
SOP-PCR-1.3	Detection and quantification of bird specific Helicobacter GFD genetic marker with quantitative polymerase chain reaction	2345962
SOP-PCR-1.4	Detection and quantification of Ruminant specific Bacteroidetes BacR genetic marker with quantitative polymerase chain reaction	2345962
SOP-PCR-1.5	Detection and quantification of canine-specific DG3 Bacteroides genetic marker with quantitative polymerase chain reaction	2345962

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
Enterolert/QT	08/03/2022	08/03/2022 16:31	Austin Nolan	08/04/2022 16:45	Jillian G Hanley	2345951, 2345952, 2345955, 2345960
SM 10200 H (mod.)	08/03/2022	08/04/2022 12:15	Ethan Wickline	08/10/2022 09:45	Carina A. Tull	2345945, 2345946, 2345948, 2345949
SM 9222 D	08/03/2022	08/03/2022 16:30	Jillian G Hanley	08/04/2022 14:53	Jillian G Hanley	2345953, 2345954, 2345957
	08/03/2022	08/03/2022 16:40	Jillian G Hanley	08/04/2022 14:53	Jillian G Hanley	2345961
SM 9223 B Quanti-Tray	08/03/2022	08/03/2022 16:41	Austin Nolan	08/04/2022 11:12	Austin Nolan	2345950, 2345959
SOP-PCR-1.0	08/03/2022	08/04/2022 10:20	Claudia Castro	08/29/2022 14:25	DeAsia Armster	2345962
SOP-PCR-1.2	08/03/2022	08/04/2022 10:20	Claudia Castro	09/01/2022 10:48	Roberta Tatti	2345962
SOP-PCR-1.3	08/03/2022	08/04/2022 10:20	Claudia Castro	09/02/2022 09:19	DeAsia Armster	2345962
SOP-PCR-1.4	08/03/2022	08/04/2022 10:20	Claudia Castro	08/30/2022 09:34	DeAsia Armster	2345962
SOP-PCR-1.5	08/03/2022	08/04/2022 10:20	Claudia Castro	08/31/2022 09:46	Roberta Tatti	2345962