

Biological Analysis Report

NE-DIST-2019-08-21-01

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

Overflow Analyses Performed By:
Advanced Environmental Laboratories - Gainesville
6815 SW Archer Road
Gainesville, Florida 32608
DOH Accreditation E82001

Event Description: **Alachua County August SMP**
Request ID: **RQ-2019-08-05-90**
Customer: **NE-DIST**
Project ID: **SMP**

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Date Certified: 01-OCT-2019 12:39



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: Sweetwater Branch - N of SE 4th St.

Collection Date/Time: 08/20/2019 11:45

Field ID: G1NE0916

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2113056	SM 10200 H (mod.)	Chlorophyll-a, Corrected	0.82	U	ug/L	P369849	
		Phaeophytin-a	0.90	U	ug/L	P369849	
		Chlorophyll-a, Uncorrected	0.60	U	ug/L	P369849	
2113061	EPA 1603	Escherichia coli-Membrane Filter	430		cfu/100 mL		
2113066	SOP-PCR-1.0	HF183-qPCR**	94	U	GEU/100 mL	P369163	
	SOP-PCR-1.2	GULL2-qPCR**	5.6E+03	U	TSC/100 mL	P369163	
	SOP-PCR-1.3	GFD-purified-qPCR**	5.31E+03	J	TSC/100 mL	P370609	MS
	SOP-PCR-1.4	BacR-qPCR**	1.0E+03	U	TSC/100 mL	P369163	
	SOP-PCR-1.5	DG3-qPCR**	860	U	TSC/100 mL	P369163	

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.

SOP-PCR-1.3: Quality control failure(s) observed. See QA report for details.

Sample Location: Sweetwater Branch - N of SE 4th St.

Collection Date/Time: 08/20/2019 11:45

Field ID: G1NE0916DUP

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2113057	SM 10200 H (mod.)	Chlorophyll-a, Corrected	0.82	U	ug/L	P369849	
		Phaeophytin-a	0.90	U	ug/L	P369849	
		Chlorophyll-a, Uncorrected	0.60	U	ug/L	P369849	
2113062	EPA 1603	Escherichia coli-Membrane Filter	180		cfu/100 mL		
2113067	SOP-PCR-1.0	HF183-qPCR**	94	I	GEU/100 mL	P369163	
	SOP-PCR-1.4	BacR-qPCR**	760	U	TSC/100 mL	P369163	
	SOP-PCR-1.5	DG3-qPCR**	640	U	TSC/100 mL	P369163	

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: Sweetwater Branch - Upstream of SR 331

Collection Date/Time: 08/20/2019 12:40

Field ID: G1NE0905

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2113058	SM 10200 H (mod.)	Chlorophyll-a, Corrected	1.6	I	ug/L	P369849	
		Phaeophytin-a	0.90	U	ug/L	P369849	
		Chlorophyll-a, Uncorrected	1.7	I	ug/L	P369849	
2113063	EPA 1603	Escherichia coli-Membrane Filter	390		cfu/100 mL		
2113068	SOP-PCR-1.0	HF183-qPCR**	110	I	GEU/100 mL	P369163	
	SOP-PCR-1.4	BacR-qPCR**	1.0E+03	U	TSC/100 mL	P369163	
	SOP-PCR-1.5	DG3-qPCR**	860	U	TSC/100 mL	P369163	

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: Bivens Arm Lake - west of US 441, and south of SW 16th Street

Collection Date/Time: 08/20/2019 14:00

Field ID: G1NE0904

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
2113059	SM 10200 H (mod.)	Chlorophyll-a, Corrected	65		ug/L	P369849	
		Phaeophytin-a	13		ug/L	P369849	
		Chlorophyll-a, Uncorrected	75		ug/L	P369849	
2113064	EPA 1603	Escherichia coli-Membrane Filter	16	B	cfu/100 mL		
2113069	SOP-PCR-1.0	HF183-purified-qPCR**	51	IJ	GEU/100 mL	P370609	LCS, SUR
	SOP-PCR-1.4	BacR-purified-qPCR**	2.0E+03	J	TSC/100 mL	P370609	LCS, SUR
	SOP-PCR-1.5	DG3-purified-qPCR**	4.9E+03	J	TSC/100 mL	P370609	LCS, SUR

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.

SOP-PCR-1.0: Quality control failure(s) observed. See QA report for details. Precision data unavailable for Analysis Batch A94433

SOP-PCR-1.4: Quality control failure(s) observed. See QA report for details. Precision data unavailable for Analysis batch A94445

SOP-PCR-1.5: Quality control failure(s) observed. See QA report for details. Precision data unavailable for Analysis batch A94457

Quality Assurance Report

Method Blank Results

Reference Method: SM 10200 H (mod.)

Batch ID: P369849

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: P369163

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

Reference Method: SOP-PCR-1.0

Batch ID: P370609

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

Reference Method: SOP-PCR-1.2

Batch ID: P369163

Component	Result	Code	Units
GULL2-qPCR	1.7E+04	U	TSC/100 mL

Reference Method: SOP-PCR-1.3

Batch ID: P370609

Component	Result	Code	Units
GFD-purified-qPCR	340	U	TSC/100 mL

Reference Method: SOP-PCR-1.4

Batch ID: P369163

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

Quality Assurance Report Method Blank Results

Reference Method: SOP-PCR-1.4
Batch ID: P370609

Component	Result	Code	Units
BacR-purified-qPCR	510	U	TSC/100 mL

Reference Method: SOP-PCR-1.5
Batch ID: P369163

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

Reference Method: SOP-PCR-1.5
Batch ID: P370609

Component	Result	Code	Units
DG3-purified-qPCR	430	U	TSC/100 mL

Quality Assurance Report Laboratory Control Sample Accuracy

Reference Method: SM 10200 H (mod.)
Batch ID: P369849

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
Chlorophyll-a, Corrected	87.6		P	84 - 116

Reference Method: SOP-PCR-1.0
Batch ID: P369163

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-qPCR	133	150	P/P	50 - 175

Reference Method: SOP-PCR-1.0
Batch ID: P370609

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
HF183-purified-qPCR	145	16.0	P/F	50 - 175

Reference Method: SOP-PCR-1.2
Batch ID: P369163

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
GULL2-qPCR	185	167	P/P	50 - 300

Reference Method: SOP-PCR-1.3
Batch ID: P370609

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
GFD-purified-qPCR	166	107	P/P	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P369163

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: P369163

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
BacR-qPCR	89.9	72.7	P/P	50 - 175

Reference Method: SOP-PCR-1.4
Batch ID: P370609

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
BacR-purified-qPCR	140	38.6	P/F	50 - 175

Reference Method: SOP-PCR-1.5
Batch ID: P369163

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
DG3-qPCR	92.2	71.6	P/P	50 - 175

Reference Method: SOP-PCR-1.5
Batch ID: P370609

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
DG3-purified-qPCR	137	47.3	P/F	50 - 175

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.0
Batch ID: P369163

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2112421	HF183-qPCR	119	128	P/P	50 - 175
2112429	HF183-qPCR	109		P	50 - 175
2112732	HF183-qPCR	107		P	50 - 175
2112733	HF183-qPCR	94.1		P	50 - 175
2112870	HF183-qPCR	96.2		P	50 - 175
2113066	HF183-qPCR	109		P	50 - 175
2113067	HF183-qPCR	113		P	50 - 175
2113068	HF183-qPCR	112		P	50 - 175
2113227	HF183-qPCR	106		P	50 - 175
2113228	HF183-qPCR	103		P	50 - 175
2113229	HF183-qPCR	98.0		P	50 - 175
2113559	HF183-qPCR	87.7		P	50 - 175

Reference Method: SOP-PCR-1.0
Batch ID: P370609

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2113069	HF183-purified-qPCR	160		P	50 - 175

Reference Method: SOP-PCR-1.2
Batch ID: P369163

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2112421	GULL2-qPCR	110	103	P/P	50 - 300

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.2

Batch ID: P369163

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2113066	GULL2-qPCR	105		P	50 - 300

Reference Method: SOP-PCR-1.3

Batch ID: P370609

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2112421	GFD-purified-qPCR	545	505	F/F	50 - 175
2113066	GFD-purified-qPCR	475		F	50 - 175
2114584	GFD-purified-qPCR	459		F	50 - 175
2114849	GFD-purified-qPCR	519		F	50 - 175
2116097	GFD-purified-qPCR	559		F	50 - 175
2116098	GFD-purified-qPCR	444		F	50 - 175
2116099	GFD-purified-qPCR	423		F	50 - 175

Reference Method: SOP-PCR-1.4

Batch ID: P369163

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2112421	BacR-qPCR	79.6	80.8	P/P	50 - 175
2112429	BacR-qPCR	53.9		P	50 - 175
2113066	BacR-qPCR	64.9		P	50 - 175
2113067	BacR-qPCR	61.5		P	50 - 175
2113068	BacR-qPCR	60.5		P	50 - 175

Reference Method: SOP-PCR-1.4

Batch ID: P370609

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2113069	BacR-purified-qPCR	130		P	50 - 175

Reference Method: SOP-PCR-1.5

Batch ID: P369163

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2112421	DG3-qPCR	92.2	90.4	P/P	50 - 175
2113066	DG3-qPCR	86.2		P	50 - 175
2113067	DG3-qPCR	88.0		P	50 - 175
2113068	DG3-qPCR	84.6		P	50 - 175

Reference Method: SOP-PCR-1.5

Batch ID: P370609

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
2113069	DG3-purified-qPCR	104		P	50 - 175

Quality Assurance Report Precision

Quality Assurance Report Precision

Reference Method: SM 10200 H (mod.)
Batch ID: P369849

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2113136	Chlorophyll-a, Corrected	2.06	Sample	P	0 - 25
2113136	Chlorophyll-a, Uncorrected	1.80	Sample	P	0 - 25

Reference Method: SOP-PCR-1.0
Batch ID: P369163

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2112421	HF183-qPCR	7.40	Spike	P	0 - 25

Reference Method: SOP-PCR-1.2
Batch ID: P369163

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2112421	GULL2-qPCR	6.87	Spike	P	0 - 25

Reference Method: SOP-PCR-1.3
Batch ID: P370609

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2112421	GFD-purified-qPCR	7.27	Spike	P	0 - 25

Reference Method: SOP-PCR-1.4
Batch ID: P369163

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2112421	BacR-qPCR	1.52	Spike	P	0 - 25

Reference Method: SOP-PCR-1.5
Batch ID: P369163

Replicated Lab Sample	Component	% RSD/RPD	Sample/Spike/LCS*	Pass/Fail	Control Limits
2112421	DG3-qPCR	1.99	Spike	P	0 - 25

* 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: 2113066
Field Sample ID: G1NE0916

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	95.4	P	50 - 300
SOP-PCR-1.2	SKETA-qPCR	126	P	50 - 300

Quality Assurance Report Surrogates

Lab Sample ID: 2113066
 Field Sample ID: G1NE0916

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.4	SKETA-qPCR	92.9	P	50 - 300
SOP-PCR-1.5	SKETA-qPCR	104	P	50 - 300

Lab Sample ID: 2113067
 Field Sample ID: G1NE0916DUP

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	91.0	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	87.0	P	50 - 300
SOP-PCR-1.5	SKETA-qPCR	103	P	50 - 300

Lab Sample ID: 2113068
 Field Sample ID: G1NE0905

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	94.2	P	50 - 300
SOP-PCR-1.4	SKETA-qPCR	90.2	P	50 - 300
SOP-PCR-1.5	SKETA-qPCR	93.7	P	50 - 300

Lab Sample ID: 2113069
 Field Sample ID: G1NE0904

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.0	SKETA-qPCR	12.9	F	50 - 300
SOP-PCR-1.4	SKETA-qPCR	11.1	F	30 - 300
SOP-PCR-1.5	SKETA-qPCR	7.39	F	50 - 300

Quality Assurance Report Summary

Ref. Method	Analyte	LCS % Recovery		MS % Recovery			Precision		
							LCS	SMP	MS
SM 10200 H (mod.)	Chlorophyll-a, Corrected	87.6						2.06	
	Chlorophyll-a, Uncorrected							1.80	
SOP-PCR-1.0	HF183-purified-qPCR	16.0	145	160					
	HF183-qPCR	150	133	119	128	109			7.40
				107					
SOP-PCR-1.2	GULL2-qPCR	167	185	110	103	105			6.87
SOP-PCR-1.3	GFD-purified-qPCR	107	166	545	505	475			7.27
				559					
SOP-PCR-1.4	BacR-purified-qPCR	38.6	140	130					
	BacR-qPCR	72.7	89.9	79.6	80.8	53.9			1.52
				64.9					
SOP-PCR-1.5	DG3-purified-qPCR	47.3	137	104					
	DG3-qPCR	71.6	92.2	92.2	90.4	86.2			1.99
				88.0					

Reference Method Descriptions

Method / DoH Cert #	Description	Associated Samples
EPA 1603 / E82001	E. coli by MF by AEL in Gainesville	2113061, 2113062, 2113063, 2113064

Reference Method Descriptions

Method / DoH Cert #	Description	Associated Samples
SM 10200 H (mod.) / E31780	Phytoplankton chlorophyll-a corrected, uncorrected, and phaeophytin by spectrophotometry	2113056, 2113057, 2113058, 2113059
SOP-PCR-1.0 / E31780	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	2113066, 2113067, 2113068
SOP-PCR-1.0 / E31780	Detection and quantification of human-specific HF183 Bacteroides genetic marker with quantitative polymerase chain reaction	2113069
SOP-PCR-1.2 / E31780	Detection and quantification of coastal bird specific Catellicoccus marimammalium Gull2 genetic marker with quantitative polymerase chain reaction	2113066
SOP-PCR-1.3 / E31780	Detection and quantification of bird specific Helicobacter GFD genetic marker with quantitative polymerase chain reaction	2113066
SOP-PCR-1.4 / E31780	Detection and quantification of Ruminant specific Bacteroidetes BacR genetic marker with quantitative polymerase chain reaction	2113066, 2113067, 2113068
SOP-PCR-1.4 / E31780	Detection and quantification of Ruminant specific Bacteroidetes BacR genetic marker with quantitative polymerase chain reaction	2113069
SOP-PCR-1.5 / E31780	Detection and quantification of canine-specific DG3 Bacteroides genetic marker with quantitative polymerase chain reaction	2113066, 2113067, 2113068
SOP-PCR-1.5 / E31780	Detection and quantification of canine-specific DG3 Bacteroides genetic marker with quantitative polymerase chain reaction	2113069

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
EPA 1603	08/20/2019	08/20/2019 15:20		08/21/2019 16:05	AEL - Gainesville	2113061, 2113062, 2113063, 2113064
SM 10200 H (mod.)	08/21/2019	08/21/2019 14:01	Yuliya Danyuk	08/28/2019 09:14	Yuliya Danyuk	2113056, 2113057, 2113058, 2113059
SOP-PCR-1.0	08/21/2019	08/21/2019 14:00	Angela Smith	09/12/2019 14:03	Mailin Lopez	2113066, 2113067, 2113068
	08/21/2019	08/21/2019 14:00	Mailin Lopez	09/23/2019 13:10	Mailin Lopez	2113069
SOP-PCR-1.2	08/21/2019	08/21/2019 14:00	Angela Smith	09/19/2019 10:48	Mailin Lopez	2113066
SOP-PCR-1.3	08/21/2019	08/21/2019 14:00	Mailin Lopez	09/23/2019 09:32	Mailin Lopez	2113066
SOP-PCR-1.4	08/21/2019	08/21/2019 14:00	Angela Smith	09/12/2019 12:51	Mailin Lopez	2113066, 2113067, 2113068
	08/21/2019	08/21/2019 14:00	Mailin Lopez	09/23/2019 14:03	Mailin Lopez	2113069
SOP-PCR-1.5	08/21/2019	08/21/2019 14:00	Angela Smith	09/13/2019 15:43	Mailin Lopez	2113066, 2113067, 2113068
	08/21/2019	08/21/2019 14:00	Mailin Lopez	09/23/2019 10:23	Mailin Lopez	2113069