

Biological Analysis Report

NW-DIST-2016-10-21-01

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

Event Description: **Bayou Chico BMAP**
Request ID: **RQ-2016-09-19-15**
Customer: **NW-DIST**
Project ID: **BMAP**

Send Reports to:
FL Dept. of Environmental Protection
160 Government Center
Suite 308
Pensacola, FL 32502-5794
Attn: Cheryl Bunch

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Certified by: Jacqueline Savage, Environmental Specialist III

Date Certified: 30-NOV-2016 16:39

A handwritten signature in black ink that reads "Jackie Savage". The signature is written in a cursive, flowing 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 group are included in this report: Microbiology.

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 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: Pensacola Bay at Sanders Beach

Collection Date/Time: 09/19/2016 10:10

Field ID: A

Matrix: W-SURF-SLT

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1843013	SOP-PCR-1.2	GULL2-qPCR**	1490	T	TSC/100 mL	P313624	

Ref. Method and Comment:
SOP-PCR-1.2: Precision data not available.

Sample Location: Bayou Chico at Mandalay Park

Collection Date/Time: 09/19/2016 10:35

Field ID: B

Matrix: W-SURF-SLT

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1843014	SOP-PCR-1.2	GULL2-qPCR**	893	T	TSC/100 mL	P313624	

Ref. Method and Comment:
SOP-PCR-1.2: Precision data not available.

Sample Location: Jones Creek at Old Corry Road

Collection Date/Time: 09/19/2016 11:11

Field ID: C

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1843015	SOP-PCR-1.2	GULL2-qPCR**	2.8E+04	U	TSC/100 mL	P313624	

Ref. Method and Comment:
SOP-PCR-1.2: No amplification of GULL2 marker. Precision data not available.

Sample Location: Jackson Creek at Old Corry Road

Collection Date/Time: 09/19/2016 11:40

Field ID: D

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1843016	SOP-PCR-1.2	GULL2-qPCR**	2.1E+04	U	TSC/100 mL	P313624	

Ref. Method and Comment:
SOP-PCR-1.2: No amplification of GULL2 marker. Precision data not available.

Sample Location: Jackson Creek at New Warrington Road

Collection Date/Time: 09/19/2016 12:10

Field ID: E

Matrix: W-SURF-FRH

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1843017	SOP-PCR-1.2	GULL2-qPCR**	2.8E+04	U	TSC/100 mL	P313624	

Ref. Method and Comment:
SOP-PCR-1.2: No amplification of GULL2 marker. Precision data not available.

Sample Location: Bayou Chico at Navy Blvd Bridge

Collection Date/Time: 09/19/2016 13:30

Field ID: F

Matrix: W-SURF-SLT

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1843018	SOP-PCR-1.2	GULL2-qPCR**	2.8E+04	U	TSC/100 mL	P313624	

Ref. Method and Comment:
SOP-PCR-1.2: No amplification of GULL2 marker. Precision data not available.

Sample Location: Bayou Chico at W Street

Collection Date/Time: 09/19/2016 13:50

Field ID: G

Matrix: W-SURF-SLT

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
1843019	SOP-PCR-1.2	GULL2-qPCR**	2.8E+04	U	TSC/100 mL	P313624	

Field ID: G

Matrix: W-SURF-SLT

Sample ID	Ref. Method	Component	Result	Code	Units	Batch ID	QC Codes
Ref. Method and Comment: SOP-PCR-1.2: No amplification of GULL2 marker. Precision data not available.							

Quality Assurance Report Method Blank Results

Reference Method: SOP-PCR-1.2
 Batch ID: P313624

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

Quality Assurance Report Laboratory Control Sample Accuracy

Reference Method: SOP-PCR-1.2
 Batch ID: P313624

Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
GULL2-qPCR	77.0		P	50 - 300

Quality Assurance Report Matrix Spike Accuracy

Reference Method: SOP-PCR-1.2
 Batch ID: P313624

Spiked Sample	Component	% Rec.1	% Rec.2	Pass/Fail	Control Limits
1843013	GULL2-qPCR	94.8		P	50 - 300
1843014	GULL2-qPCR	104		P	50 - 300
1843015	GULL2-qPCR	63.0		P	50 - 300
1843016	GULL2-qPCR	88.4		P	50 - 300
1843017	GULL2-qPCR	94.4		P	50 - 300
1843018	GULL2-qPCR	84.5		P	50 - 300
1843019	GULL2-qPCR	101		P	50 - 300

Quality Assurance Report Surrogates

Lab Sample ID: 1843013
 Field Sample ID: A

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.2	SKETA-qPCR	111	P	50 - 300

Lab Sample ID: 1843014
 Field Sample ID: B

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.2	SKETA-qPCR	110	P	50 - 300

Quality Assurance Report Surrogates

Lab Sample ID: 1843015
 Field Sample ID: C

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.2	SKETA-qPCR	108	P	50 - 300

Lab Sample ID: 1843016
 Field Sample ID: D

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.2	SKETA-qPCR	121	P	50 - 300

Lab Sample ID: 1843017
 Field Sample ID: E

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.2	SKETA-qPCR	124	P	50 - 300

Lab Sample ID: 1843018
 Field Sample ID: F

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

Lab Sample ID: 1843019
 Field Sample ID: G

Reference Method	Surrogate	% Rec.	Pass/Fail	Control Limits
SOP-PCR-1.2	SKETA-qPCR	119	P	50 - 300

Quality Assurance Report Summary

Ref. Method	Analyte	LCS % Recovery	MS % Recovery	LCS	Precision SMP	MS
SOP-PCR-1.2	GULL2-qPCR	77.0	94.8 104 63.0 88.4			

Reference Method Descriptions

Method / DoH Cert #	Description	Associated Samples
SOP-PCR-1.2 / E31780	Detection and quantification of coastal bird specific <i>Catellibacoccus marimammali</i> genetic marker with quantitative polymerase chain reaction	1843013, 1843014, 1843015, 1843016, 1843017, 1843018, 1843019

Preparation and Analysis Log

Ref. Method	Received Date	Prep Date/Time	Prepared By	Analysis Date/Time	Analyzed By	Associated Samples
SOP-PCR-1.2	09/20/2016	09/20/2016 13:00	Sarah Menz	10/20/2016	Sarah Menz	1843013, 1843014, 1843015, 1843016, 1843017, 1843018, 1843019