

eDNA Reporting Template

I. eDNA Testing Sample Submission Information				
Report Title:				
Project Number:		Date of Final Reporting:		
Service Provider Information	Type:		Requesting Organization Information	
	Contact Name:			Organization Name:
	Address:			Contact Name:
	Contact Phone:			Contact Phone:
	Contact Email:			Contact Email:
LAB ACCREDITATION / CERTIFICATION:				
Executive Summary - Study Objectives, Rationale, and Main Finding(s) derived from both eDNA samples and controls				
Appendices (Required)		Check to confirm inclusion	Appendices (Additional)	
Appendix 1: Maps of the study sites and sampling locations		<input type="checkbox"/>	Appendix 5:	
Appendix 2: Contamination prevention procedures		<input type="checkbox"/>	Appendix 6:	
Appendix 3: qPCR protocol		<input type="checkbox"/>	Appendix 7:	
Appendix 4: Metadata and qPCR data		<input type="checkbox"/>	Appendix 8:	
II. Study Design and eDNA Sampling				
A. Study information	A.1 Species targeted (common and Latin):			
	A.2 Study objectives:			
	A.3 Geographic location and/or region:			
	A.4 Sampling date (range):	Start:	Finish:	
	A.5 Sample types:			
	A.6 Mapping databases (list all):			
B. Study design	B.1 Type(s) of ecosystem:			
	B.2 Sampling design (how does sampling optimize species detection for study goal?):			
	B.3 Number of sites sampled:			
	B.4 Number of stations sampled within sites (add explanation for variation among sites):			
	B.5 Number of field sample replicates:			
	B.6 Time series (number of times sites and stations were sampled):			
	B.7 Environmental conditions, relevant observations, and additional field data:			
	B.8 Field blanks and field controls (describe and give numbers):			
C. eDNA sample collection	C.1 Env. sample collection method:			
	C.2 Volume / weight sampled:			
	C.3 Sample depth(s):			
	C.4 Field sample storage/time to processing:			
	C.5 Sample processing method (list disposable equipment; preservative used):			
	C.6 Filter type and pore size:			
	C.7 Sample preservation:			

III. eDNA Sample Analysis - Laboratory Methods		
D. DNA extraction	<i>D.1 Name of commercial kit or protocol:</i>	
	<i>D.2 Reference protocol:</i>	
	<i>D.3 DNA extraction controls:</i>	
	<i>D.4 Proportion of total sample:</i>	
	<i>D.5 DNA elution volume:</i>	
	<i>D.6 Extracted eDNA storage conditions:</i>	
E. qPCR assay	<i>E.1 Assay Name:</i>	
	<i>E.2 Assay Type:</i>	
	<i>E.3 Level of assay validation:</i>	
	<i>E.4 Specificity data:</i>	
	<i>E.5 Dilution and volume of DNA used:</i>	
	<i>E.6 qPCR positive and negative controls:</i>	
	<i>E.7 Technical replicates per sample:</i>	
	<i>E.8 Inhibition tests:</i>	
	<i>E.9 Number of qPCR cycles:</i>	
IV. Summary of eDNA Results		
F. Reporting control results	<i>F.1 Criteria to determine if controls passed or failed:</i>	
	<i>F.2 Positive control results (report each type separately):</i>	
	<i>F.3 Negative control results (report each type separately):</i>	
	<i>F.4 Failed controls (report and explain):</i>	
G. Reporting eDNA sample results	<i>G.1 Calculated LOD:</i>	
	<i>G.2 QA/QC qPCR results:</i>	
	<i>G.3 Other qPCR results:</i>	
	<i>G.4 Determination of sample-level results:</i>	
	<i>G.5 Determination of station-level results:</i>	
	<i>G.6 Determination of site-level results:</i>	
H. Closing statements	<i>H.1 Disclaimer (any additional information to help explain results for any samples, stations, or sites):</i>	
	<i>H.2 Summary of eDNA detection:</i>	
	<i>H.3 Future recommendations:</i>	