

### Helbing Laboratory eDNA Technical Bulletin

All eDNA tools are validated through a rigorous multi-step evaluation protocol that includes tests of DNA target specificity and amplification sensitivity<sup>1-3</sup>.

#### General eDNA Assay Information

Target Species: White sturgeon (*Acipenser transmontanus*)      eDNA qPCR Tool: eACTR4      Gene Target: MT-Dloop  
 Species Code: te-ACTR      eDNA qPCR Format: TaqMan      Published in:

#### eDNA Assay Sensitivity Test Summary using gBlocks™ Synthetic DNA

LOD 0.4      95% CI 0.3-0.8 Copies/Rxn      LOQ 1.5      95% CI 1-2.8 Copies/Rxn      LOB 0 hits/8  
 LOQ<sub>continuous</sub> 4 Copies/Rxn

Binomial-Poisson model for 8 technical replicates determined using eLowQuant R code<sup>4</sup>.      When the LOQ < LOD, use the LOD for the LOQ.

Enzyme: QIAcuity

#### eDNA Assay Specificity Test Information

Each qPCR reaction in the specificity assay contained 10 picograms of voucher target gDNA (n=25 technical replicates)

Species	Common Name ( <i>Species</i> )	Detection	# Voucher Specimens	Sample Sources/Locations
ma-CALUfa	Domestic dog ( <i>Canis lupus familiaris</i> )	No	1	British Columbia
ma-FECA	Domestic cat ( <i>Felis catus</i> )	No	1	British Columbia
ma-HOSA	Human ( <i>Homo sapiens</i> )	No	1	Netherlands
te-ACFU	Lake sturgeon/yellow sturgeon ( <i>Acipenser fulvescens</i> )	Yes*	2	Quebec
te-ACME	Green sturgeon ( <i>Acipenser medirostris</i> )	No	3	California
te-ACTR	White sturgeon ( <i>Acipenser transmontanus</i> )	Yes	15	British Columbia (Upper and Lower Fraser River, Nechako River, Columbia River, Kootenay Lake)
te-CAAU	Goldfish ( <i>Carassius auratus</i> )	No	1	British Columbia
te-CACA	Longnose sucker ( <i>Catostomus catostomus</i> )	No	1	British Columbia
te-CACAch	Salish sucker ( <i>Catostomus catostomus (chehalis)</i> )	No	1	British Columbia
te-CACO	White sucker ( <i>Catostomus commersonii</i> )	No	1	Ontario
te-CAMA	Largescale sucker ( <i>Catostomus macrocheilus</i> )	No	1	Ontario
te-CLPA	Pacific herring ( <i>Clupea pallasii</i> )	No	1	British Columbia
te-COAR	Cisco/Tubilee ( <i>Coregonus artedii</i> )	No	1	Alberta
te-COCL	Lake whitefish ( <i>Coregonus clupeaformis</i> )	No	1	Alberta
te-COCO	Slimy sculpin ( <i>Cottus cognatus</i> )	No	1	British Columbia
te-ESLU	Northern pike ( <i>Esox lucius</i> )	No	1	British Columbia
te-GAAC	Threespine stickleback ( <i>Gasterosteus aculeatus</i> )	No	1	British Columbia
te-HYPR	Hypomesus pretiosus ( <i>Hypomesus pretiosus</i> )	No	1	British Columbia
te-LEGI	Pumpkinseed ( <i>Lepomis gibbosus</i> )	No	1	British Columbia
te-LOLO	Burbot ( <i>Lota lota</i> )	No	1	Alberta
te-MIDO	Smallmouth bass ( <i>Micropterus dolomieu</i> )	No	1	British Columbia
te-MISA	Peamouth chub ( <i>Mylocheilus caurinus</i> )	No	1	British Columbia
te-ONCLle	Westslope cutthroat trout ( <i>Oncorhynchus clarkii lewisi</i> )	No	1	Alberta
te-ONGO	Pink salmon ( <i>Oncorhynchus gorbuscha</i> )	No	1	British Columbia
te-ONKE	Chum salmon ( <i>Oncorhynchus keta</i> )	No	1	British Columbia
te-ONKI	Coho salmon ( <i>Oncorhynchus kisutch</i> )	No	1	British Columbia
te-ONMY	Rainbow (steelhead) trout ( <i>Oncorhynchus mykiss</i> )	No	1	Alberta
te-ONNE	Sockeye salmon ( <i>Oncorhynchus nerka</i> )	No	1	British Columbia
te-ONTS	Chinook salmon ( <i>Oncorhynchus tshawytscha</i> )	No	1	British Columbia
te-PEFL	Yellow perch ( <i>Perca flavescens</i> )	No	1	Quebec
te-PRWI	Mountain whitefish ( <i>Prosopium williamsoni</i> )	No	1	Alberta
te-PTOR	Northern pikeminnow ( <i>Ptychocheilus oregonensis</i> )	No	1	Ontario
te-RHCA	Longnose (nooksack) dace ( <i>Rhinichthys cataractae</i> )	No	1	Ontario
te-SACO	Bull trout ( <i>Salvelinus confluentus</i> )	No	1	Alberta
te-SAFO	Brook trout ( <i>Salvelinus fontinalis</i> )	No	1	Alberta
te-SAMA	Dolly varden ( <i>Salvelinus malma</i> )	No	1	British Columbia
te-SANA	Lake trout ( <i>Salvelinus namaycush</i> )	No	1	Alberta
te-SPTH	Longfin smelt ( <i>Spirinchus thaleichthys</i> )	No	1	Washington
te-THPA	Oolichan/Eulachon ( <i>Thaleichthys pacificus</i> )	No	1	British Columbia

\*1/25 replicates obtained a positive hit

#### References

- Hobbs, J, Adams, IT, Round, JM, Goldberg, CS, Allison, MJ, Bergman, LC, Mirabzadeh, A, Allen, H, Helbing, CC (2020) Revising the range of Rocky Mountain tailed frog, *Ascaphus montanus*, in British Columbia, Canada, using environmental DNA methods. Environmental DNA, 2: 350-361. <https://doi.org/10.1002/edn3.82>
- Hobbs, J, Round, JM, Allison, MJ, Helbing, CC (2019) Expansion of the known distribution of the coastal tailed frog, *Ascaphus truei*, in British Columbia, Canada, using robust eDNA detection methods. PLOS ONE 14(3): e0213849. <https://doi.org/10.1371/journal.pone.0213849>
- Langlois, VS, Allison, MJ, Bergman, LC, To, TA, and Helbing, CC (2020) The need for robust qPCR-based eDNA detection assays in environmental monitoring and risk assessments. Environmental DNA, 3: 519-527. doi: 10.1002/edn3.164
- Lesperance, M, Allison, MJ, Bergman, LC, Hocking, MD, and Helbing, CC (2021) A statistical model for calibration and computation of detection and quantification limits for low copy number environmental DNA samples. Environmental DNA, 3: 970-981. doi: 10.1002/edn3.220

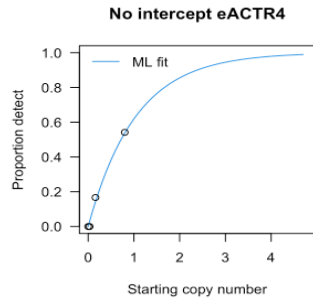
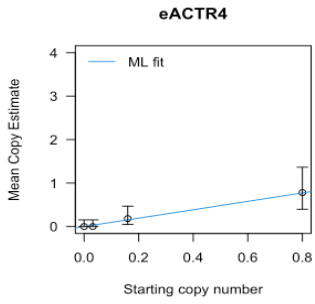
eDNA Assay Sensitivity Test Details using gBlocks™ synthetic DNA

To calculate tables for different numbers of replicates, raw csv data files can be accessed here:  
<https://onlineacademiccommunity.uvic.ca/helbinglab/edna/>

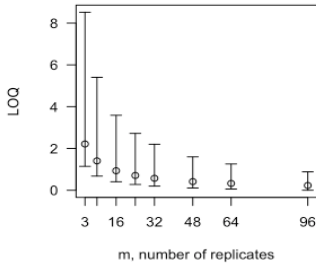
From 8 Technical Replicates

# Detects	# Copies	SE
0	0	0
1	0.138	0.143
2	0.298	0.224
3	0.487	0.308
4	0.718	0.407
5	1.016	0.536
6	1.437	0.727
7	2.155	1.106

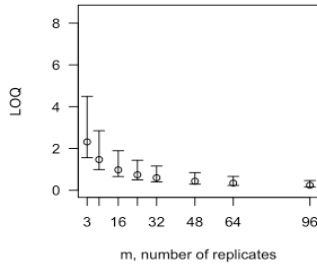
Determined using eLowQuant R code<sup>4</sup>.



Limits quant - intercept eACTR4

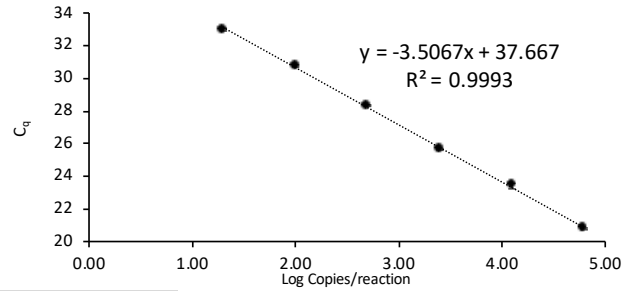


Limits quant - no intercept eACTR4



Binomial-Poisson model: No intercept  
Determined using eLowQuant R code<sup>4</sup>.  
Based on a 2 µL DNA input in a total 15 µL reaction

Applied to reactions with ≥95% positive hits



Efficiency 93%

Field Sample Validation

Sample Type	Known		Detected	Location
	Presence	# Samples		

Abbreviations

95% CI	95% Confidence interval	LOQ	Limit of quantification
eDNA	Environmental DNA	MT-Dloop	Mitochondrial displacement loop (D-loop) gene
gDNA	Total genomic DNA extracted from voucher specimen	NTC	qPCR no template control
LOB	Limit of blank	qPCR	Quantitative real-time polymerase chain reaction
LOD	Limit of detection	SE	Standard error