

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 sensitivity¹⁻³.

General eDNA Assay Information

Target Species: Oregon Spotted Frog (*Rana pretiosa*) eDNA qPCR Tool: eRAPR2 Gene Target: MT-CYB
 Species Code: am-RAPR 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 1-2.9 Copies/Rxn LOB 0 hits/8
 LOQ_{continuous} 4 Copies/Rxn

Binomial-Poisson model for 8 technical replicates determined using eLowQuant R code⁴. 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		Sample Sources/Locations
			Specimens		
am-ANBO	Western Toad (<i>Anaxyrus (Bufo) boreas</i>)	No	1		Northwest Territories
am-ASMO	Rocky Mountain Tailed Frog (<i>Ascaphus montanus</i>)	No	1		British Columbia
am-LICA	Bullfrog (<i>Lithobates (Rana) catesbeiana</i>)	No	1		British Columbia
am-PSRE	Pacific Chorus Frog (<i>Pseudacris (Hyla) regilla</i>)	No	1		British Columbia
am-RAAU	Northern Red-legged Frog (<i>Rana aurora</i>)	No	1		British Columbia
am-RACA	Cascades Frog (<i>Rana cascadae</i>)	No	1		British Columbia
am-RALU	Columbia Spotted Frog (<i>Rana luteiventris</i>)	Yes*	1		British Columbia
am-RAPR	Oregon Spotted Frog (<i>Rana pretiosa</i>)	Yes	1		British Columbia
am-TAGR	Rough-skinned Newt (<i>Taricha granulosa</i>)	No	1		British Columbia
ma-HOSA	Human (<i>Homo sapiens</i>)	No	1		Netherlands

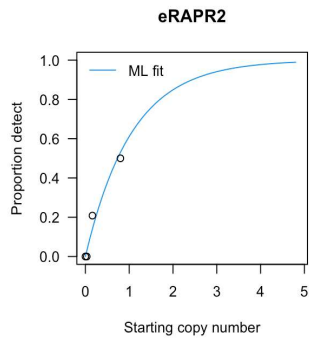
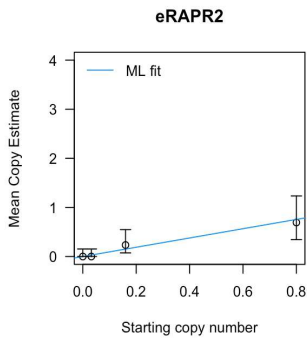
References

- 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

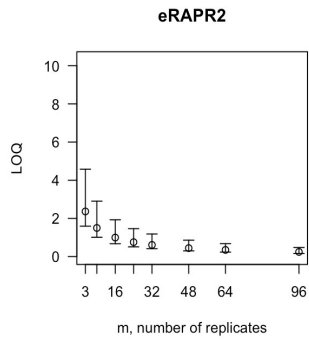
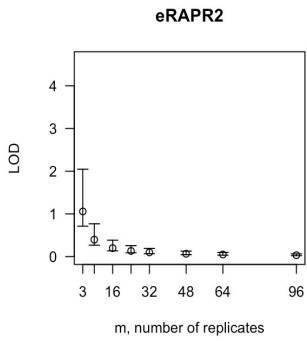
To calculate tables for different numbers of replicates, raw csv data files can be accessed here:
<https://onlineacademiccommunity.uvic.ca/helbinglab/edna/>

From 8 Technical Replicates

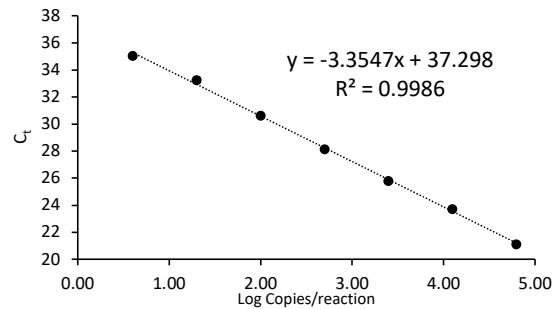


# Detects	# Copies	SE
0	0	0
1	0.14	0.15
2	0.31	0.23
3	0.5	0.32
4	0.73	0.42
5	1.04	0.55
6	1.47	0.74
7	2.2	1.13

Determined using eLowQuant R code⁴.



Applied to reactions with $\geq 95\%$ positive hits



Efficiency 99%

Binomial-Poisson model: No intercept
 Determined using eLowQuant R code⁴.
 Based on a 2 μ L DNA input in a total 15 μ L reaction

Field Sample Validation

Sample Type	Known		Detected	Location
	Presence	# Samples		

Abbreviations

95% CI	95% Confidence interval	LOQ	Limit of quantification
eDNA	Environmental DNA	MT-CYB	Mitochondrial cytochrome B 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