

Supplementary information for

‘Fishing in the water: effect of sampled water volume on environmental DNA-based detection of macroinvertebrates’

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Appendix I PCR Conditions

Figure S1 eDNA concentrations relative to used sample volumes

Appendix I: PCR conditions

PCR mixture

Our PCR mixtures contained 1 X Buffer (Roche, Switzerland), 0.18 mM dNTP, 1 X BSA, 0.05 U/μL Taq (Roche, Switzerland), 2.0 mM MgCl₂ and 0.5 μM of each primer (see table below) and 2 μL of eDNA in a total reaction volume of 15 μL.

Primer

Polymerase chain reaction (PCR) primer pair sequences and annealing temperatures. Amplicon size includes primer pair lengths.

Species	Primer name	Primer sequence 5'–3'	Annealing temperature (°C)
<i>Ancylus fluviatilis</i>	Anf-F2	AATAGTTGAAGGAGGAGCAG	56
	Anf-R2	CAAATAAGGAAAGACGTTCC	
<i>Baetis buceratus</i>	Bab-F	GACTGCTACCCCTTCATTG	51
	Bab-R	GCATGCGATCCAATGTTATC	
<i>Gammarus pulex</i>	Gap-F	TTTTGACTTTTACCTCCTTCYC	48.5
	Gap-R	GATATACCAGGTCTACGTATATTG	

PCR temperature regime

50 cycles

95° C	4 min
95° C	30 sec
Annealing temperature dependent on used primer, see table above	30 sec
72° C	1 min
72° C	5 min
10° C	forever

Figure S1: eDNA concentrations relative to used sampled volumes

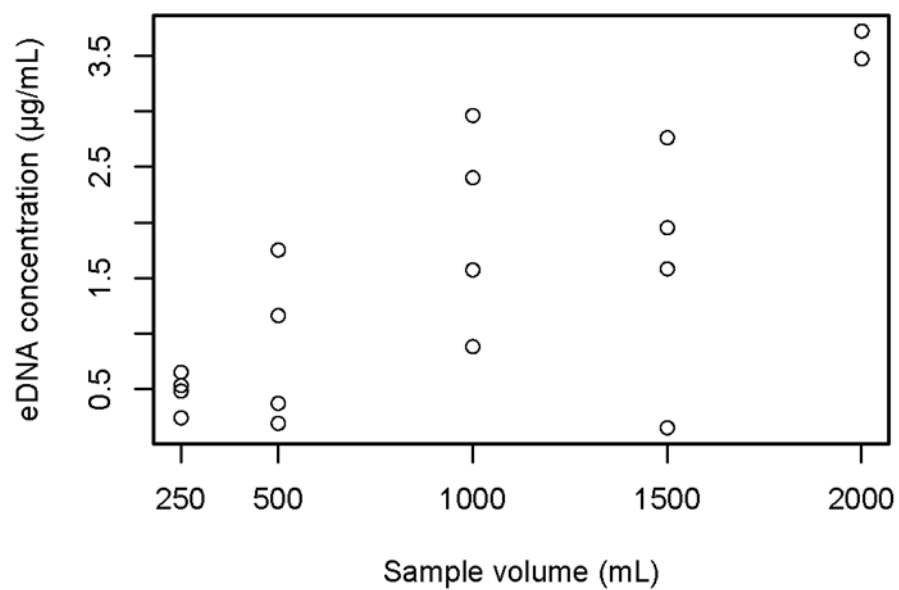


Fig. S1: eDNA concentration relative to the used sample volume. Two data points are missing for 2000 mL as we run out of sample to measure DNA concentration.