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[Modified] Lake ABPS Protocol - University of Maine

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ABSTRACT

A modified version of the Lake ABPS protocol as described in Thomson-Laing et al. 2022

Protocol successful at detecting fish sedDNA collected from lake **surface** sediments, as well as river sediments during an anadromous fish sea-run migration

OPEN ACCESS

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Protocol status: Working
We use this protocol and it's working

Created: Mar 09, 2023




















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PROTOCOL integer ID:
78446


Keywords: Sedimentary DNA, SedDNA, Fish


Alkaline Lysis & Ethanol Precipitation


4h 1m

- 1 **CENTRIFUGE** sediment samples at  5250 x g for  00:05:00 5m
DISCARD pore water using a sterile pipette, so only sediment remains
- 2 **ADD**  10 g of wet sediment to a sterile 50 mL tube
ADD  6 mL sodium hydroxide (NaOH, 0.33M) to the sample
ADD  3 mL Tris-EDTA (pH 8.0) to the sample
- 3 **VORTEX** sample at max speed for  00:01:00 56m
INCUBATE sample at  65 °C for  00:55:00
ALLOW samples to cool to  Room temperature
- 4 **CENTRIFUGE** samples at  5250 x g for  01:00:00 1h
TRANSFER  7.5 mL of supernatant to a new, sterile 50 mL tube
- 5 **ADD**  7.5 mL Tris-HCl (pH 6.7)
ADD  1.5 mL sodium acetate (3M, pH 5.2)
ADD  30 mL molecular grade ethanol
- 6 **INCUBATE** samples at  -20 °C for  01:00:00 1h
- 7 **CENTRIFUGE** sample at  5250 x g for  01:00:00 1h
DISCARD supernatant
ALLOW remaining ethanol to evaporate off of concentrated pellet before proceeding to next step

PowerSoil Pro Extraction on Concentrated Pellet - sample p.. 29m

8 **WEIGH** the concentrated pellet and split it into multiple  0.5 g replicates

TRANSFER each  0.5 g replicate into a PowerBead Pro Tube


9 **ADD**  800 µL of Solution CD1 to each PowerBead Pro Tube



20m

SECURE PowerBead Pro Tubes horizontally to a Vortex Adapter

VORTEX for  00:10:00

ROTATE tubes so caps are oriented in the opposite direction

VORTEX for another  00:10:00

10 **CENTRIFUGE** sample at  15000 x g for  00:02:00

2m

TRANSFER all supernatant to a clean 2 mL Microcentrifuge Tube

PowerSoil Pro Extraction on Concentrated Pellet - inhibitor ..

29m

11 **ADD**  200 µL of Solution CD2

VORTEX briefly to mix

12 **CENTRIFUGE** at  15000 x g for  00:01:00

1m


AVOIDING the pellet, transfer **all** supernatant to a clean 2 mL Microcentrifuge Tube

PowerSoil Pro Extraction on Concentrated Pellet - bind DNA

29m

13 **ADD**  600 µL of Solution CD3

VORTEX briefly to mix

14 **LOAD**  650 µL of lysate onto an MB spin column

1m


CENTRIFUGE at  15000 x g for  00:01:00

DISCARD the liquid flow-through

15 REPEAT step 14 to ensure all the lysate has passed through the MB Spin Column


CAREFULLY place the MB spin column into a clean 2mL collection tube

PowerSoil Pro Extraction on Concentrated Pellet - wash spi... 29m

16 ADD  500 μ L of Solution EA to the MB spin column 1m



CENTRIFUGE at  15000 x g for  00:01:00

DISCARD the liquid flow-through and place the MB spin column into the same 2 mL Collection Tube


17 ADD  500 μ L of Solution C5 to the MB spin column 1m

CENTRIFUGE at  15000 x g for  00:01:00

DISCARD the liquid flow-through and place the MB Spin Column into a **new** 2 mL Collection Tube

18 CENTRIFUGE at  16000 x g for  00:02:00 2m

CAREFULLY place the MB spin column into a **new** 2mL Collection Tube



19 ADD  50-100 μ L of Solution C6 to the center of the white membrane in the MB Spin Column

Note

Adjust the amount of Solution C6 added to each replicate so that the final volume, once all replicates are pooled together (step 21), totals 200ul

For example, if at Step 8 sample A weighed 1.0g and was split into two 0.5g replicates: A1 and A2. At this step (Step 19), A1 and A2 would each receive 100ul Solution C6, so that when they are pooled together, their total volume is 200ul

If sample A was split into three 0.5g replicates (A1, A2, and A3), each would receive approximately 66ul of Solution C6

20 **CENTRIFUGE** at  15000 x g for  00:01:00


1m

DISCARD the MB Spin Column

POOL all replicates into a sterile 1.5 mL Microcentrifuge Tube

DNA is now ready for downstream applications

Note

For best results in qPCR, use ~  6 µL of extracted DNA template per PCR reaction