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CTAB Extraction Protocol for Sediment

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Supplementary Appendix A1.

CTAB Extraction Protocols for Sediment and Water
Extraction protocols were modified from Cocco et al. (2005, 2006, 2007).

Sediment Extraction Protocol (steps involving centrifugation should be performed inside a fume hood)

1. Thaw the CTAB preserved sediment sample in the fridge for no more than 24 hours.
2. Once thawed, accurately measure the content of the 10-ml tube with 10% bleach and rinse with reverse osmosis water.
3. Vortex at highest speed for 30 sec, then incubate at 60°C for 10 min.
4. Add 15 ml of 50% Chloroform/Isoamyl alcohol (24:1).
5. Vortex the sediment/CTAB/water mixture briefly and shake at low speed (Vortexer setting 4) for 5 min.
6. Centrifuge at 3250g for 15 min at room temperature to separate aqueous and organic phases.
7. Without touching the intermediate layer, carefully transfer the aqueous phase (supernatant) to a new 15-ml tube (Tip: Use a 10-ml serological pipette for the first 8 to 12 ml, then a 1000 µl micropipette to aspirate the last 2 to 3 ml.)
8. Add an equal volume of 100% isopropanol and 1 volume of 5M NaCl to the supernatant and chill in a -20°C freezer for 1 hr (or overnight if more convenient).
9. Centrifuge at 3250g for 15 min at room temperature, carefully pour off the supernatant.
10. Add 1 ml of 70% EtOH, washing down the inner walls of the tube, then centrifuge at 3250g for 2 min at room temperature.
11. Pour off EtOH and allow the DNA pellet to air dry completely (use a 45°C incubator to expedite ethanol evaporation).
12. Resuspend the pellet in 100 µl of LoTE buffer. Heat briefly at 45°C and swirl gently to mix and resuspend. Once fully resuspended, briefly centrifuge to collect all liquid in the bottom of 15-ml tube.
13. Transfer all liquid to a 1.5-ml, low-bind microcentrifuge tube.
14. Use 200 µl in QIAquick® Spin Purification Kit (Qiagen Research, Irvine, CA). This can now be used in genetic assays.

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Protocol status: Working

Successfully used by Turner et al. (2015) to detect carp sedDNA

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Abstract

Successfully used by Turner et al., 2015 to detect bigheaded Asian carp surface sedimentary DNA from experimental ponds and natural rivers

<https://www.sciencedirect.com/science/article/pii/S000632071400442X>

Materials


Materials

[M] 100 millimolar (mM) Tris-HCL 

[M] 1.4 Molarity (M) NaCl


[M] 1 Mass / % volume Polyvinylpyrrolidone 

[M] 2 Mass / % volume Cetyl trimethyl ammonium bromide (CTAB)

[M] 20 millimolar (mM) EDTA 

Troubleshooting

Safety warnings

 Steps involving Sevag should be performed inside a fume hood.



Sample preparation

1d 1h 47m 30s

- 1 **THAW** the CTAB-preserved sediment sample in the fridge for no more than

24:00:00

1d

DECONTAMINATE the exterior of the sample tube with 10 % bleach solution and rinse with reverse osmosis water

- 2 **VORTEX** at max speed for 00:00:30

10m 30s

INCUBATE at 60 °C for 00:10:00

Chloroform extraction

1d 1h 47m 30s

- 3 **ADD** 15 mL of Sevag (Chloroform/Isoamyl alcohol 24:1)

Note

Steps involving Sevag should be performed inside a fume hood.

- 4 **VORTEX** the sediment/CTAB/Sevag mixture briefly

5m

SHAKE at low speed (Vortexer setting 4) for 00:05:00

- 5 **CENTRIFUGE** at 3220 x g for 00:15:00 at Room temperature

15m

CAREFULLY transfer the supernatant to a new 50 mL tube

Ethanol precipitation

1d 1h 47m 30s

- 6 **ADD** an equal volume of ice-cold Isopropanol to the supernatant

1h

ADD ½ volume of 5M NaCl to the supernatant


INCUBATE at -20 °C for 01:00:00 (or overnight if more convenient)

- 7 **CENTRIFUGE** at 3220 x g for 00:15:00 at Room temperature

15m


CAREFULLY pour off the supernatant



8 **ADD**  2 mL of 70% EtOH, washing down the inner walls of the tube



2m

CENTRIFUGE at  3220 x g for  00:02:00 at  Room temperature

POUR off EtOH and allow the DNA pellet to air dry completely (a  45 °C incubator can be used to evaporate stubborn droplets)

DNA resuspension

1d 1h 47m 30s

9 **RESUSPEND** the pellet in  1000 µL of LoTE buffer. Heat briefly at  45 °C and swirl gently to mix and resuspend. Once fully resuspended, briefly centrifuge to collect all liquid in the bottom of 50-mL tube.

10 Transfer all liquid to a 1.5-µL low-bind microcentrifuge tube

11 Use  200 µL in OneStep™ Inhibitor Removal Kit (Zymo Research, Irvine, CA)

Protocol references

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