



Version 2 ▼

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Quick Fungal DNA extraction from colonies on plates V.2

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1 Works for me



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ABSTRACT

Adapted from Chi, M. H., Park, S. Y., & Lee, Y. H. (2009).

Colony PCR on fungal colonies grown on plates does not work as well as it does for bacteria (it usually doesn't work at all). DNA therefore needs to be extracted first. As this DNA will only be used as a PCR template to check for presence / absence of individual genes I am not too concerend about high molecular weight or purity. This protocol is quick and can be used to process several samples at the same time.

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PROTOCOL CITATION

Version created by Johannes Wolfram Debler

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Make stock solutions

1

1M TrisHCl pH 8.0	100 ml
12.14 g	Tris base or Trizma base
up to 100 ml	deionised H ₂ O
Adjust pH	concentrated HCl

2M KCI (Potassium chloride)	200 ml
15.11 g	KCI (Potassium chloride)
up to 200 ml	deionised H ₂ O

Α	В
500 mM NaEDTA (pH 8.0)	100 ml
18.6 g	EDTA disodium salt dihydrate
up to 100 ml	deionised H ₂ O

Adjust pH to 8.0, otherwise the EDTA won't go into solution

Autoclave stock solutions at 121°C

Make Extraction Buffer

Extraction 100 Buffer ml 2M KCl 50 ml 1M Tris-HCl 10 pH 8.0 ml 500 mM 2 ml NaEDTA deionised up H_2O to 100

Preparation

3 Per sample prepare 2 x 1.5 ml Eppendorf tubes. Label them with sample number and add:

Tube 1:500 ul Extraction Buffer + 1 stainless steel bead

Tube 2: 300 ul Isopropanol

Harvest fungal material

With a **sterile toothpick or pipette tip** remove a fungal colony or a piece of mycelia of about 3-5 mm x 3-5 mm in size from a plate and put it into **Tube 1** (containing the Extraction Buffer and steel bead).

Try not to use a piece that is too big, as that might lead to carryover of too many PCR inhibitors. I usually use my toothpick like a sewing machine needle, punching out a piece of mycelium all the way to the bottom of the agar plate. Then I use the toothpick to slice underneath the actual mycelial mat to try to cut off as much of the agar as possible.

Homogenise fungal material

5 Homogenise samples for 5 min at 1500 rpm in the MiniG 1600.

SPEX SP 1600

MiniG 1600 Automated Tissue Homogenizer and Cell Lyser



Pellet Cell Debris

6 Pellet cell debris by centrifugation at 17.000 x g for 5 min.

DNA Precipitation

- 7 Transfer supernatant to **Tube 2** (Containing 300 ul Isopropanol)
 - Vortex for 5 10 seconds
 - Pellet DNA by centrifugation for 5 min at 17.000 x g

Wash DNA

- 8 Discard supernatant
 - Add 800 ul 70% Ethanol
 - Vortex mix sample, make sure pellet comes loose from bottom of the tube
 - Pellet DNA by centrifugation for 5 min at 17.000 x g

Dissolve DNA

- 9 Discard supernatant
 - Remove as much of the ethanol as possible by drawing drops out with a pipette tip
 - Let leftover ethanol evaporate for 5-10 min
 - Dissolve pellet in 50-100 ul H₂O or 1x TE buffer
 - Measure DNA concentration (e.g. Nanodrop)
 - store DNA at -20°C