

Aug 10, 2023

# 🌐 Labyrinthulomycete total RNA extraction protocol - hot phenol

DOI

[dx.doi.org/10.17504/protocols.io.q26g7pyo8gwz/v1](https://dx.doi.org/10.17504/protocols.io.q26g7pyo8gwz/v1)



Jackie Collier<sup>1</sup>

<sup>1</sup>Stony Brook University

Protist Research to Opti...

Collier Lab



Jackie Collier

Stony Brook University

## Create & collaborate more with a free account

Edit and publish protocols, collaborate in communities, share insights through comments, and track progress with run records.

Create free account

OPEN  ACCESS



DOI: <https://dx.doi.org/10.17504/protocols.io.q26g7pyo8gwz/v1>

**Protocol Citation:** Jackie Collier 2023. Labyrinthulomycete total RNA extraction protocol - hot phenol. **protocols.io**  
<https://dx.doi.org/10.17504/protocols.io.q26g7pyo8gwz/v1>



### Manuscript citation:

Lippmeier, J.C., Crawford, K.S., Owen, C.B., Rivas, A.A., Metz, J.G. and Apt, K.E. (2009), Characterization of Both Polyunsaturated Fatty Acid Biosynthetic Pathways in *Schizochytrium* sp.. Lipids, 44: 621-630. <https://doi.org/10.1007/s11745-009-3311-9>

**License:** This is an open access protocol distributed under the terms of the **Creative Commons Attribution License**, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited

**Protocol status:** Working

**We use this protocol and it's working**

**Created:** August 10, 2023

**Last Modified:** August 10, 2023

**Protocol Integer ID:** 86355

**Keywords:** labyrinthulomycete total rna extraction protocol, part of the labyrinthulomycete jgi community sequencing project, labyrinthulomycete jgi community sequencing project, labyrinthulomycete, total rna extraction protocol, rna extraction, rna, hot phenol modified, sequencing

## Abstract

Modified from Lippmeier et al. 2009; developed as part of the labyrinthulomycete JGI Community Sequencing Project and Gordon and Betty Moore Foundation Marine Microbial Eukaryote Transcriptome Sequencing Project (MMETSP)

## Troubleshooting

## Safety warnings

❗ Be super careful with hot phenol! It's near the boiling point of phenol, so just a little too hot can be explosive.



## Preparing biomass and reagents

- 1 Grow up cells, collect, and freeze rapidly - preferably in liquid nitrogen. Store biomass at -80C if not extracting immediately.

This protocol has worked so far for two different thraustochytrids (*Aurantiochytrium limacinum* ATCC MYA-1381, *Schizochytrium aggregatum* ATCC 28920) and two *Aplanochytrium* strains (PBS06 and PBS07).

- 2 Prepare extraction buffer

100 mM Tris-HCl pH 7.5

1.5 M NaCl

50 mM Na<sub>2</sub>-EDTA pH 8.0

20 g per liter CTAB (cetyltrimethylammonium bromide)

8 mg per ml DTT (dithiothreitol) - ADD FRESH JUST BEFORE USE

(note, if you keep these stocks in glass bottles, first soak the bottles with 0.1 M NaOH to inactivate RNases)

Prepare Tris-equilibrated phenol, pH 4.5-4.8, by warming to 65 C

Prepare acid phenol:chloroform:isoamyl alcohol (pH 4.5-4.8, 125:24:1) by warming to 65 C

Set up to incubate the extraction step at 65 C

Prepare 8M LiCl

Chill 100% ethanol and 70% ethanol

Prepare TE or nuclease-free water

Get a bucket of ice

## Extraction steps - REPEAT 3 TIMES (do 3 extractions)

- 3 Suspend ~1000 micrograms wet weight biomass per ~10 ml extraction buffer by vortexing

(these amounts would yield tens of micrograms of total RNA)

- 3.1 Incubate at 65 C for 5 minutes, mix by vortex or inverting 2 or 3 times

- 4 Add equal volume hot (65 C) acid phenol/chloroform/isoamyl alcohol, vortex to mix well



- 4.1 Incubate at 65 C for 5 minutes, mix by vortex or inverting 2 or 3 times
- 5 Cool the mixture in ice
- 6 Centrifuge to separate the phases 4 C, 6,000g, 15 minutes
- 7 Move the top phase to a new tube, avoiding the interface

## Precipitation steps

- 8 Add an equal volume of isopropanol and incubate overnight at 4 C
- 8.1 If remaining DNA forms a large fluffy precipitate, remove by spooling onto glass rod
- 9 Collect precipitate by centrifugation 4 C, 16,000g, 30 minutes  
(may require distributing the sample among several 1.5 ml microfuge tubes and/or multiple spins for each tube to process the volume)
- 10 Wash each pellet with 0.5 ml ice-cold 75% ethanol twice, centrifuging 4 C 16,000g, 5 minutes
- 10.1 Allow the pellets to air dry
- 11 Dissolve pellets in an appropriate volume, analyze by appropriate method (spectrophotometry, Bioanalyzer); store frozen at -80 C

## Optional cleanup steps

- 12 If the RNA contains particulates that do not dissolve, they can be removed by centrifugation 4 C, 6,000g, 5 minutes and transferring the supernatant to a fresh tube.



- 13 Even if the RNA does not have obvious DNA, a DNase treatment step is nonetheless recommended
- 14 If the RNA contains large amounts of DNA or other contaminants, it can be removed by selectively precipitating the RNA with 17 microliters of 8M LiCl per 50 microliters RNA
  - 14.1 Incubate -20 C 60 minutes
  - 14.2 Collect RNA by centrifugation 4 C, 16,000g, 20 minutes
- 15 Wash each pellet with 0.5 ml ice-cold 75% ethanol twice, centrifuging 4 C 16,000g, 5 minutes
  - 15.1 Allow the pellets to air dry
- 16 Dissolve pellets in an appropriate volume, analyze by appropriate method (spectrophotometry, Bioanalyzer); store frozen at -80 C