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NOV 07, 2022



WORKS FOR ME 1

Cyanobacteria Total Lipid Extraction from Cell Pellets
Forked from Cyanobacteria Total Lipid Extraction from Polycarbonate Filters

DOI

dx.doi.org/10.17504/protocols.io.4r3l2oe3jv1y/v1

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The Aquatic Microbial Ecology Research Group - AMERG (The Buchan, Zinser and Wilhelm labs) CyanoHABs <u>1 more workspa</u>



ABSTRACT

This protocol is designed/used for extraction of total cellular lipids from cyanobacteria samples (either lab cultures or field samples) collected via centrifugation for use in lipid analysis and quantification *via* mass spectrometry.

Please contact Dr. Steven Wilhelm (wilhelm@utk.edu) or Robbie M. Martin (rmarti49@vols.utk.edu) for additional information regarding this protocol.

Modified from Guan, X. L., Riezman, I., Wenk, M. R., & Riezman, H. (2010). Yeast lipid analysis and quantification by mass spectrometry. *Methods in Enzymology*, *470*, 369-391.

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PROTOCOL CITATION

Robbie Martin, Steven W Wilhelm, Katarina A. Jones, Shawn Campagna 2022. Cyanobacteria Total Lipid Extraction from Cell Pellets. **protocols.io**

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 $MANUSCRIPT\ CITATION\ please\ remember\ to\ cite\ the\ following\ publication\ along\ with\ this\ protocol$

Guan, X. L., Riezman, I., Wenk, M. R., & Riezman, H. (2010). Yeast lipid analysis and quantification by mass spectrometry. Methods in Enzymology, 47 369-391.

FORK NOTE



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FORK FROM

Forked from Cyanobacteria Total Lipid Extraction from Polycarbonate Filters, Robbie Martin

KEYWORDS

cyanobacteria, lipids, extraction

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CREATED

May 12, 2022

LAST MODIFIED

Nov 07, 2022

PROTOCOL INTEGER ID

62491

- 1 Prepare the three separate solutions needed for this extraction protocol as follows:
 - lipid extraction solvent: a 15:15:5:1:0.18 ratio by volume of 95% ethanol, water, diethyl ether, pyridine, and 4.2 N ammonium hydroxide, respectively.
 - water-saturated butanol: a 1:1 ratio of butanol and Milli-Q water
 - purified lab water: (Milli-Q water)
- 2 Concentrate cells via centrifugation using methods appropriate for your available lab equipment and for your species of interest.

 Decant supernatant. Transfer pelleted cells to a 2-mL centrifuge tube. Cells may need to be re-pelleted and residual supernatant removed after transfer to final 2-mL centrifuge tube.

Note: Appropriate volume of lab culture or field samples to concentrate and extract depends on cell concentration. As a guideline, we have been successful concentrating 10-25 mL of lab cultures of *Microcystis aeruginosa* and ~50 mL of either raw lake water or mesocosm samples.

- 3 Add 1 mL of extraction solvent, \sim 100 μ L of glass beads, and vortex \sim 5 s.
- 4 Incubate sample in 60 °C water bath for 20 min.
- 5 Centrifuge sample at 10,000 x g for 10 min.
- Remove supernatant and place into a 1-dram glass vial (dram vial #1). The first two extractions from a sample will be placed in this vial (#1).
- 7 Repeat steps 3-6, except DO NOT ADD more glass beads.



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8	Dry the collected supernatant in dram vial #1 under a stream of nitrogen.
9	Re-suspend dried sample in 300 μL of water-saturated butanol and 150 μL of Milli-Q water.
10	Vortex and transfer to a 2-mL centrifuge tube.
11	Centrifuge at 10,000 x g for 2 min.
12	Remove top butanol phase and place into a NEW 1-dram glass vial (dram vial #2).
13	Wash original dram vial (#1) with 300 µL saturated butanol and transfer to residual aqueous phase in 2-mL centrifuge tube from step 10. Vortex.
14	Centrifuge at 10,000 x g for 2 min. Remove top butanol phase and place into dram vial #2.
15	Dry the collected butanol phase in dram vial #2 under a stream of nitrogen.
16	Re-suspend dried sample in 300 μ L of 9:1 methanol:chloroform.
17	The samples are now ready for analysis <i>via</i> LC/MS.