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Cyanobacteria Total Lipid Extraction from Polycarbonate Filters

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DOI

dx.doi.org/10.17504/protocols.io.byxtpxnn[Robbie Martin](#)¹, [Steven W Wilhelm](#)¹, Katarina A. Jones¹, [Hector Castro](#)¹, [Shawn Campagna](#)¹¹University of Tennessee, Knoxville[The Aquatic Microbial Ecology Research Group - AMERG \(The Buchan, Zinser and Wilhelm labs\)](#)[CyanoHABS](#)[1 more workspace](#)

Robbie Martin

COMMENTS 0

ABSTRACT

This protocol is designed/used for extraction of total cellular lipids from cyanobacteria samples (either lab cultures or field samples) collected on polycarbonate filters 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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KEYWORDS

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- 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 Unfold polycarbonate filter and place into a 2-mL centrifuge tube with cell side of filter facing outwards.

Note: Appropriate volume of lab culture or field samples to filter and extract depends on cell concentration. As a guideline, we have been successful filtering 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, ~100 µL of glass beads, and vortex ~5 s.
- 4 Incubate sample in 60 °C water bath for 20 min.
- 5 Centrifuge sample at 10,000 x g for 10 min.
- 6 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.

- 8 Dry the collected supernatant in dram vial #1 under a stream of nitrogen.
- 9 Re-suspend dried sample in 300 μ L of water-saturated butanol and 150 μ L 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 μ L 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 *via* LC/MS.