

Dec 17, 2024

Isolation of Lipid Droplets from Cultured Cortical Neurons

 In 1 collection

DOI

<https://dx.doi.org/10.17504/protocols.io.n2bvj934plk5/v1>

Mukesh Kumar¹, Daehun Park^{2,3,4}, Pietro De Camilli^{2,3,4}, Timothy A. Ryan^{1,2}

¹Department of Biochemistry, Weill Cornell Medicine, New York, NY 10065;

²Aligning Science Across Parkinson's (ASAP) Collaborative Research Network, Chevy Chase, Maryland 20815, USA;

³Department of Neuroscience, Howard Hughes Medical Institute, Program in Cellular Neuroscience, Neurodegeneration and Repair, Yale University School of Medicine, New Haven, Connecticut 06520, USA;

⁴Department of Cell Biology, Yale University School of Medicine, New Haven, Connecticut 06520, USA



Mukesh Kumar

Weill Cornell Medical College

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.n2bvj934plk5/v1>

Protocol Citation: Mukesh Kumar, Daehun Park, Pietro De Camilli, Timothy A. Ryan 2024. Isolation of Lipid Droplets from Cultured Cortical Neurons. **protocols.io** <https://dx.doi.org/10.17504/protocols.io.n2bvj934plk5/v1>

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: November 26, 2024

Last Modified: December 17, 2024

Protocol Integer ID: 114539

Keywords: ASAPCRN, lipid droplets from cultured cortical neuron, purifying lipid droplet, isolation of lipid droplet, lipid droplet, cultured cortical neurons this protocol, cultured cortical neuron, lipidomic analysis, constituent lipid, suitable for lipidomic analysis, neuron

Funders Acknowledgements:

NIH

Grant ID: NS036942

NIH

Grant ID: NS11739

ASAP

Grant ID: ASAP-024404

Abstract

This protocol provides a robust method for purifying lipid droplets from cultured cortical neurons, suitable for lipidomic analysis of constituent lipids.

Guidelines

The protocol needs prior approval by the users' Institutional Animal Care and Use Committee (IACUC) or equivalent ethics committee.



Materials

- Cortical neurons: 8-10 × 15 cm dishes (~80% confluent) 14-21 DIV
-  KLMH45 Merck MilliporeSigma (Sigma-Aldrich) Catalog #SML1998
- MEPS Buffer Components:
 1.  PIPES Merck MilliporeSigma (Sigma-Aldrich) Catalog #P6757
 2.  EGTA Merck MilliporeSigma (Sigma-Aldrich) Catalog #E3889
 3.  Magnesium sulfate solution Merck MilliporeSigma (Sigma-Aldrich) Catalog #M3409
 4.  Potassium hydroxide ACS reagent, ≥85%, pellets Merck MilliporeSigma (Sigma-Aldrich) Catalog #221473-500G
- cOmplete Protease Inhibitor Cocktail (Roche, 45-11697498001)
-  Sucrose Merck MilliporeSigma (Sigma-Aldrich) Catalog #S0389
-  BODIPY[®] 493/503 (4,4-Difluoro-1,3,5,7,8-Pentamethyl-4-Bora-3a,4a-Diaza-s-Indacene) ThermoFisher Catalog #D3922
-  CellMask[®] Orange Plasma membrane Stain ThermoFisher Catalog #C10045

Equipment:

- Nitrogen Cavitation Apparatus
- Centrifugation Equipment:
 1. Table-top centrifuge (capacity 2 ml, up to 1800 g),
 2. Ultracentrifuge with swinging-bucket rotor (SW41Ti, Beckman Coulter; up to 1,74,000 g)
- Needles and Syringes: 25 gauge needles with 2 ml syringes.



Neuronal Preparation

- 1 Isolate cortical neurons from P0-P1 pups and culture them in 15 cm dishes (8-10 plates).
- 2 Maintain the neurons in standard culture condition for 14-21 days to allow growth of axons and synapse formation.
- 3 Treat the neurons with **5 micromolar (μM)** KLH45 (dissolved in DMSO) for 24 hours prior to harvesting. 

Harvesting and Lysis

- 4 Harvest neurons in **1.5 mL** of **0.9 Mass Percent** MEPS buffer supplemented with 1X protease inhibitor cocktail. 

MEPS buffer composition: PIPES: 35 mM (adjust pH to 7.2 with KOH), EGTA: 5 mM, MgSO_4 : 5 mM, Sucrose: 0.9 M

- 5 Lyse the neurons using the nitrogen cavitation method or Isobiotec Cell Homogenizer (8-10 strokes with 16-micron clearance). Confirm complete lysis under an inverted white-field or phase-contrast microscope.

Post-Nuclear Supernatant (PNS) Preparation

10m

- 6 Centrifuge the lysate at **1800 x g, 4°C, 00:10:00** to remove unlysed somas, nuclei, and plasma membrane fractions.  

10m

- 7 Collect the supernatant (called post-nuclear supernatant, PNS) into a fresh tube.

Gradient Setup and Ultracentrifugation

1h 30m

- 8 Pour the PNS over 2.5 M MEPS in a clear ultracentrifuge tube (SW41).



9 Overlay the bottom layer with  2 mL each of [M] 1.2 Mass Percent ,
[M] 0.5 Mass Percent , and [M] 0 Mass Percent MEPS to establish a step gradient.



10 Load the ultracentrifuge tube into the SW41Ti rotor and balance it against an identical tube.

11 Centrifuge the gradient at  174000 x g, 4°C, 01:30:00 .

1h 30m



Lipid Droplet (LD) Collection

12 Carefully transfer the ultracentrifuge buckets to ice.

13 Collect the LDs floating on the top of gradient using a syringe fitted with appropriate needles (25 gauge) or a tube slicer.

Storage

14 Snap-freeze the collected LDs in liquid nitrogen for future use.

Validation of Purity

15 Perform western blotting to confirm the purity of the isolated LDs.

16 Confirm enrichment of LD by Perilipin-2 and contamination (if any) using organelle-specific markers for mitochondria, ER, Golgi, and endosomes.



Note

- Appropriate clearance/number of strokes with Isobiotec cell Homogenizer, or pressure during Nitrogen cavitation is critical for effective lysis while preserving organelle integrity.
- Extreme care is required during the gradient setup to prevent contamination of the LD layer.
- Ensure all ultracentrifuge tubes are properly balanced to avoid damage of rotor/centrifuge.
- Cell-Mask Plasma Membrane Stain can be used to confirm absence of unwanted membrane contamination in LDs samples.