

Oct 17, 2024

# Sample Preparation and 3D-SIM fixed-cell imaging of FTH1 and Filipin in NPC1 and NPC2 mutants



Forked from Evaluation of pUb kinetics using 3D-SIM



In 1 collection

DOI

#### dx.doi.org/10.17504/protocols.io.3byl4qyjzvo5/v1

Felix Kraus<sup>1</sup>

<sup>1</sup>Department of Cell Biology, Blavatnik Institute, Harvard Medical School, 240 Longwood Ave, Boston MA 02115, USA

ASAP Collaborative Rese...



Felix Kraus

# OPEN ACCESS



DOI: dx.doi.org/10.17504/protocols.io.3byl4qyjzvo5/v1

**Protocol Citation:** Felix Kraus 2024. Sample Preparation and 3D-SIM fixed-cell imaging of FTH1 and Filipin in NPC1 and NPC2 mutants. **protocols.io** <u>https://dx.doi.org/10.17504/protocols.io.3byl4qyjzvo5/v1</u>

**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

Created: January 12, 2024

Last Modified: October 17, 2024

Protocol Integer ID: 93441

Keywords: ASAPCRN, Microscopy



**Funders Acknowledgement:** 

**ASAP** 

Grant ID: ASAP-000282

**ASAP** 

Grant ID: ASAP-025160

#### Disclaimer

DISCLAIMER - FOR INFORMATIONAL PURPOSES ONLY; USE AT YOUR OWN RISK

The protocol content here is for informational purposes only and does not constitute legal, medical, clinical, or safety advice, or otherwise; content added to **protocols.io** is not peer reviewed and may not have undergone a formal approval of any kind. Information presented in this protocol should not substitute for independent professional judgment, advice, diagnosis, or treatment. Any action you take or refrain from taking using or relying upon the information presented here is strictly at your own risk. You agree that neither the Company nor any of the authors, contributors, administrators, or anyone else associated with **protocols.io**, can be held responsible for your use of the information contained in or linked to this protocol or any of our Sites/Apps and Services.

#### Abstract

Protocol for the evaluation of pUb kinetics using 3D-SIM



### Seeding of HeLa cells

- 1 Wash HeLa TMEM192-3xHA control and NPC1, NPC2 mutants with 1xPBS
- 2 Add Trypsin to cells for 5 min and incubate at 37°C to dissociate cells from plastic well
- 3 Resuspend cells in 1 mL DMEM media
- 4 Count cells
- Seed appropriate number of cells onto 18x18mm Marienfeld Precision cover glasses thickness No. 1.5H (tol.  $\pm$  5  $\mu$ m).
- 6 Top up glass bottom dish with either 1 mL DMEM and place cells back into incubator
- 7 The next day exchange growth medium

## Staining

- Aspirate DMEM and fix cells in warm paraformaldehyde 3% Glutaraldehyde 0.35% in 0.1M Sodium Cacodylate, pH 7.4Aspirate PFA solution and wash wells 3x with PBST (1x PBS, 0.02% Tween 20).
  - PFA is toxic and discard in the appropriate waste streams.
- 9 Stain cholesterol with Filipin (0.05mg/ml in PBS / 10% FBS) for 2h at RT. Wash cells after staining with PBST for 2 min.
- 10 Permeabilize the cells by adding 0.2% Triton X-100 in PBS.
- 11 Remove the detergent solution by aspiration. Wash wells 3x with PBST (1x PBS, 0.02% Tween 20). Drain well.
- 12 Block cells for 10 min with 3% BSA 1x PBS.



- Remove BSA solution by aspiration. Wash wells 3x with PBST (1x PBS, 0.02% Tween 20). Drain well.
- 14 Incubate with primary antibodies in 3% BSA 1x PBS over night at 4°C with gentle shaking.
  - a. anti-FTH1 (rabbit)
- 15 Wash wells 3x with PBST (1x PBS, 0.02% Tween 20). Drain well.
- 16 Incubate with secondary antibodies in 3% BSA 1x PBS for 45 min 1h.
  - a. Goat anti-rabbit AlexaFluor 568
- 17 Wash wells 3x with PBST (1x PBS, 0.02% Tween 20). Drain well.
- 18 Wash coverslips with 1x PBS and mount in Vectashield (Vector Laboratories, H-1000-10) on glass slides. Exchange PBST with 1x PBS and keep cells at 4°C until imaging. Image within the next few days.

## Fixed-cell 3D-SIM microscopy

- Image cells on DeltaVision OMX v4 using an Olympus 60x / 1.42 Plan Apo oil objective (Olympus, Japan). Record 405, 488 and 568 channels using a front-illuminated sCMOS (PCO Photonics, USA) in 512x512px image size mode, 1x binning, 125 nm z-stepping and with 15 raw images taken per z-plane (5 phase-shifts, 3 angles).
- 20 Reconstruct raw images using CUDA-accelerated 3D-SIM reconstruction code (https://github.com/scopetools/cudasirecon) based on Gustafsson et al. (2008[FK1]). The Optimal optical transfer function (OTF) was determined via an in-house build software, developed by Talley Lambert from the NIC / CBMF (GitHub: https://github.com/tlambert03/otfsearch, all channels were registered to the 528nm output channel, Wiener filter: 0.002, background: 90).

## 3D-redendering in ChimeraX

- Open .dv composite image stacks in Fiji. Separate channels and save them as separate .tiff stacks for import into ChimeraX
- 22 Open ChimeraX and open one of the two image stacks into a new project.



- 23 While not moving the scene, drag the second tiff stack into ChimeraX. The two channels should thus keep their previous alignment.
- 24 Go Tools > Volume Data > Volume Viewer. Change the representation of both channels to surface or mesh (or what fits your visualization needs best).
  - Adjust color and levels of the image on the histogram.
- 25 In the Volume Viewer tab, use the "Hide Dust" function to remove unspecific background specs / spots from the render. Repeat for both channels.
- 26 Move the render into an adequate position and export the image.
- 27 Import the resulting image into illustrator for making your figure.