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# © TBK1 knockdown and rescue in Hela-M cells

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**ABSTRACT** 

TANK-binding kinase 1 (TBK1) is a multifunctional kinase with roles in several crucial cell processes, including innate immune response, anti-viral response, and mitochondrial clearance, the last of which is of most interest to us. In order to further understand the role of TBK1 in mitophagy, we developed a protocol to transiently deplete TBK1 from a model system, HeLa cells, and re-introduce a tagged TBK1 along with other relevant components of mitophagy. Because of the many processes that rely on TBK1, cells depleted of the kinase exhibit poor health. Our protocol accomplishes  $\sim 70\%$  depletion of endogenous TBK1 within 24 hours without causing excessive cell death. We employ this protocol to carry out biochemistry experiments such as Western blotting and organelle fractionation, and imaging experiments such as immunofluorescence, and live cell imaging.

**ATTACHMENTS** 

db5nbjnsp.pdf

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PROTOCOL CITATION

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KEYWORDS

tissue culture, transfection, knockdown, TBK1

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#### **GUIDELINES**

- This protocol was developed with the HeLa subtype, HeLa-M. HeLa-M cells are flatter than standard HeLa cells, making them easier to image. They also uptakes iRNA better than standard HeLa. Regardless, the protocol would be easily adaptable to standard HeLa cells or other cell culture lines.
- HeLa cells fare poorly after TBK1 depletion. For this reason, I developed a 24-hour protocol with simultaneous knockdown and rescue. Even so, there will be a number of dead cells in the dish after transfection.
- This protocol was created in order to investigate Parkin-dependent mitophagy. Parkin and several other fluorescently-tagged mitophagy components are included in the protocol.

#### MATERIALS TEXT

#### Materials

**⊗** Countess slides **Thermo Fisher** 

Scientific Catalog #C10228

**⊠** 10 ml conical

tube Corning Catalog #CLS430055

Sigma Catalog #EP022364120

Corporation Catalog #P35G-1.5-20-C

Reagents

Scientific Catalog #R001100

Scientific Catalog #T10282

**⊠DMEM Corning Catalog #10-017-CV** 

**⊗** FBS (HyClone) Contributed by users

⊠ GlutaMAX glucose supplement Gibco - Thermo

Fischer Catalog #35050061

**⊠** Opti-MEM Thermo Fisher

Scientific Catalog #3198507

### - Plasmid DNA

- untagged Parkin (subcloned from YFP-Parkin, a gift from R. Youle, NIH, Bethesda, MD)
- Mito-DsRed2 (kindly provided by. T. Schwartz, Harvard Medical School, Boston)
- Halo-TBK1(subcloned from SNAP-TBK1, provided by T.Maniatis, see Ye et al, PNAS, 2019)

### - siRNA

- targeting 5' (UAACAAGAGGAUUGCCUGA) end of hTBK1(Horizon Discovery)
- targeting 3' (CCACUGUUAUACUGGGAUA) end of hTBK1(Horizon Discovery)

Scientific Catalog #11668027

# Equipment

- cell incubator to maintain 37 °C and 5% CO<sub>2</sub> atmosphere
- Countess automated cell counter (Invitrogen, AMQAX1000)
- minicentrifuge (SouthernLabware, MLX-106)
- Compound microscope

### SAFETY WARNINGS

For hazard information and safety warnings, please refer to the SDS (Safety Data Sheet).

# BEFORE STARTING

- Cells are best transfected before passage 30. Higher passage number could result in lower transfection efficiency. Use cells between P5 and P25 for best results.
- siRNA is easily degraded by RNAases that exist on everyday surfaces. Follow best practices for handling siRNA by suspending desiccated reagent in RNAase-free water, aliquoting into sterilized tubes, and using barrier pipet tips to handle aliquots.

### Day 1: Plating

Trypsinize Hela-M cells by aspirating all media from a 10 cm dish of confluent cells, then dropping **Q.75 mL Trypsin** onto cells.

- Resuspend detached cells and neutralize Trypsin with 11 mL DMEM with [M]10 % FBS and [M]1 % GlutaMAX for a final volume of 11.75 mL.
- 4 Transfer this volume to 10 mL conical tube.
- 5 Combine  $\square 10 \, \mu l$  of suspension with  $\square 10 \, \mu l$  Trypan blue in a 1.5 mL tube.
- 6 Drop 10 μl of this mixture onto a Countess slide and insert into the cell counter to calculate the concentration of cells in the resuspended solution.
- 7 Plate ~0.25 million HeLaM cells on 35mm imaging dish in **2 mL DMEM/ FBS/ GlutaMAX**.

The number of cells to plate should be calculated by a standard guide for confluence, in order to achieve  $\sim$ 80-90% confluence on Day 2.

I prefer to pipet the volume of suspension needed (usually 20-50  $\mu$ L) for each dish directly from that 10 mL tube and add it dropwise to 2 mL in the dish. I found that the cells uptake the exogenous nucleic acids best when they are in confluent patches instead of evenly distributed on the surface.

# Day 2: Transfection

8 Examine cells by compound microscope **§ 18:00:00** - **§ 24:00:00** after plating to confirm 80-90% confluence.

If cells are not at 80-90% confluence, do not transfect. Wait until they reach 80-90%.

For each dish, prepare the following two solutions in 1.5 mL tubes. Tube 1 (nucleic acids): **■200 µl Opti-mem** • + **□0.5 μl Halo-TBK1** (stock at 1μg/μL) ■ + **□0.25** µl mito-dsRed (" ") • + **□**0.5 μl Parkin (" ") + ■4.8 µl siTBK1-5' + ■4.8 µl siTBK1-3' Invert stock solutions of each plasmid several times in order to ensure even distribution of plasmid. Tube 2 (Lipofectamine 2000): **200** µl Opti-mem + ■11.4 µl Lipofectamine 2000 The Lipofectamine 2000 volume is calculated by this equation: ■ (X\*4)+(Y\*3.2), where X = ug plasmid DNA (in this protocol, X=1.25) and Y = # of 4.8 µL aliquots of siRNA(in this protocol, Y = 2) Invert tubes 8 times to distribute the contents, then: 10m 11 Incubate © 00:05:00 - © 00:10:00 at § Room temperature. 2s 12 Spin © 00:00:02 in a minicentrifuge. 13 Add Tube 2 to Tube 1 and invert 8 times to mix. 15m 14 Incubate © 00:05:00 - © 00:10:00 at § Room temperature. 2s 15 mprotocols.io 08/31/2021

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Spin © 00:00:02 in a minicentrifuge.

Add entire volume (~> **400** μl) to the cells dropwise, distributing the drops mostly in the center of the dish (where the imaging window/coverslip is).

1d

Day 3: Collection 1d

17 Cells are ready to collect for various assays **§ 18:00:00** - **§ 24:00:00** after transfection step.