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🌐 Live-Imaging of GCase Activity in iPSC-Derived Dopaminergic Neurons Using the Lyso-FQ Probe

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We use this protocol and it's working

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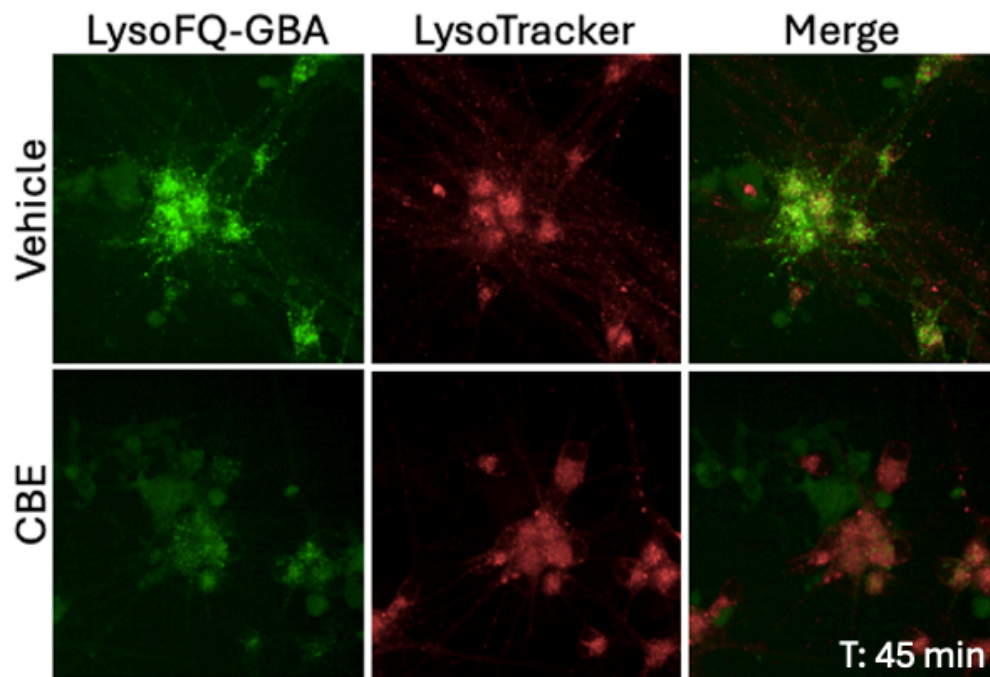
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Abstract

This protocol describes the procedure for assessing GCase activity in iPSC-derived dopaminergic neurons using the Lyso-FQ GCase substrate (Biolegend 421948). The assessment is performed via live-cell imaging with the Opera Phenix high-content imaging system and analyzed using Harmony software.

This LysoFQ-GBA (Lysosomal GCase Probe Green) probe, a cell-permeable reporter for GCase activity, fluoresces upon glucocerebrosidase activation, facilitating the detection and monitoring of GBA1 activity.



Materials

Reagents	Company	Cat. Number
LysoTracker Deep Red	ThermoFisher Scientific	L12492
Lyso-FQ	Biologend	421948
FluoroBrite DMEM	ThermoFisher Scientific	A1896701
CBE (Conduritol B epoxide)	Millipore Sigma	C5424
DMSO	FisherScientific	BP231-100

Safety warnings

- ⚠ Be extremely gentle when dispensing into the wells, as dopaminergic neurons are prone to detaching with forceful pipetting



CBE treatment

16h

- 1 For the negative control, incubate neurons overnight at $37\text{ }^{\circ}\text{C}$ with $25\text{ }\mu\text{M}$ of the GCase inhibitor Conduritol B epoxide (CBE)

16h

Labeling lysosomes with LysoTracker Deep Red

30m

- 2 Prepare a LysoTracker Deep Red working solution at 50 nM in media. Remove the existing media from the wells and add $60\text{ }\mu\text{L}$ of the LysoTracker solution. Incubate for $00:30:00$ at $37\text{ }^{\circ}\text{C}$.

30m

Preparing LysoFQ stock solution

- 3 While the cells are incubating, prepare the Lyso-FQ stock solution (Biolegend 421948). Reconstitute one $500\text{ }\mu\text{g}$ vial of Lyso-FQ in $77.3\text{ }\mu\text{L}$ of DMSO for a 5 mM stock, then aliquot into $3\text{ }\mu\text{L}$ single-use stock vials. Store aliquots at $-80\text{ }^{\circ}\text{C}$, protected from light, and avoid repeated freeze-thaw cycles.

Addition of Lyso-FQ

- 4
 1. Prepare the Lyso-FQ working solution by diluting the stock 1:1000 in FluoroBrite DMEM.
 2. Carefully remove the LysoTracker Deep Red solution from the wells and gently add $50\text{ }\mu\text{L}$ of the Lyso-FQ working solution to each well.
 3. Proceed with imaging

Imaging using the Opera Phenix high-content imaging system

2h 30m

- 5
 1. Insert the prepared plate into the Opera Phenix chamber.
 2. Utilize a $40\times$ water immersion objective. Capture images using the 488 nm and 647 nm laser channels. Acquire 15–18 fields of view per well every $00:15:00$ for a total duration of $02:30:00$. Maintain the plate under standard cell culture conditions ($37\text{ }^{\circ}\text{C}$, $5\%\text{ CO}_2$) throughout the imaging session.

2h 30m

Image Analysis Workflow in Harmony Software

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1. Load Images

Open the Data Analysis tab in the Harmony software and load the desired image dataset.

2. Create an Analysis Sequence

3. Identify Cells

Use the **Alexa 647** channel with the **P method** to detect cells. Parameters: Area > 100 μm^2 , Splitting Sensitivity: 0.70, Common Threshold: 0.55). Save this population as "**Cells**".

4. Select Cell Population

Apply Common Filters to the "Cells" population to refine it, creating a new population named "**Cells Selected**".

5. Calculate Morphological Properties

For the "**Cells Selected**" population, use the Standard Method to calculate the **Area**.

6. Calculate Intensity Properties

- For the **Alexa 647** channel, use the Standard Method to calculate the **mean intensity** for the "Cells Selected" population.
- For the **Alexa 488** channel, use the Standard Method to calculate the **mean intensity** for the "Cells Selected" population.

7. Define Results Output

- Output: **Cells Selected - Intensity Cell Alexa 647 Mean**
- Output: **Cells Selected - Intensity Cell Alexa 488 Mean**

8. Run the Analysis

9. Export Data

Download the results as a **CSV file**.

Assess GCase activity

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- From the table generated by Harmony, calculate the ratio of the PFB-FDGlu mean fluorescence intensity to the LysoTracker Deep Red (647) baseline reading (at t=0) for each time point.
- Enter these calculated values into Prism to generate the GCase activity curves.