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Measuring mitophagy via FACS with mtKeima reporter

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Protocol status: Working

We use this protocol and it's working

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Abstract

Preparation of samples for measuring mitophagy levels using mtKeima reporter by fluorescence activated cell sorting (FACS).

Troubleshooting



Day 1


- 1 Seed cells in a 24 well plate, aiming for a confluency of ~80-90% at the time of treatment. Seed additional wells of cells not expressing any fluorescent proteins, cells expressing only mtKeima, and cells expressing only YFP-Parkin (to be used as gating controls).

Day 2

2h 3m

- 2 Feed all cells with standard growth media for 01:00:00 prior to treatment. 1h
- 3 Replace media in each well with media containing the drug you are treating with.
NOTE: Do not change the media or treat the additional wells of cells to be used for gating control.
- 4 01:00:00 prior to harvesting, feed the untreated wells with 0.5 mL of fresh growth media. 1h
- 5 At the conclusion of the treatment time point, harvest the cells using the following procedure:
 - 5.1 Aspirate media from all wells.
 - 5.2 Wash all wells once with 500 μ L of Room temperature PBS.
 - 5.3 Add 150 μ L of trypsin to each well, and incubate cells at 37 $^{\circ}$ C for 00:01:30 1m 30s
 - 5.4 Place plates onto ice, and harvest each sample into a separate microfuge tube on ice by resuspending each sample with 500 μ L of ice cold standard growth media.
- 6 Centrifuge all samples at 1000 rcf for 00:01:30 at 4 $^{\circ}$ C 1m 30s
- 7 Carefully aspirate the supernatant from all samples.



- 8 Resuspend each sample in  50 μL of FACS sorting media (10% v/v FBS, 1 mM EDTA in PBS) and place into FACS analysis tubes. Keep samples on ice until immediately prior to analysis.