#### **Supplementary Information**

#### Repurposing protein degradation for optogenetic modulation of protein activities

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# **Supplementary Methods**

# Reagents

Phusion DNA polymerase master mix (M0530L) was purchased from NEB. In-Fusion HD Cloning Plus kit was from Takara (638909). DreamTaq PCR Master Mix (2×) (K1081), Turbofect (R0532) and Pierce Protease and Phosphatase Inhibitor Mini Tablets (A32959), F12K (21127-022) medium, and horse serum (26050088) were from Thermo Fisher Scientific. Fetal bovine serum (F1051), RIPA Lysis Buffer, 10× (20-188) was from Millipore Sigma. PBS (21-040-CV), DMEM (10-017-CV), Penicillin-streptomycin solution (30-002-CI), Trypsin EDTA (0.25% Trypsin, 0.1% EDTA) 1× (25-053-CI) was from Corning. Precast protein gels (456-1024) and ECL substrate (170-5060) were from Bio-Rad. Antibodies used in this work are Phospho-p44/42 MAPK (Erk1/2) (Thr202/Tyr204) (Cell Signaling Technology, 9101S, 1:1000), p44/42 MAPK (Erk1/2) (Cell Signaling Technology, #9102S, 1:1000), HA-Tag (C29F4) (Cell Signaling Technology, #3724S, 1:1000), GAPDH (14C10) (Cell Signaling Technology, 7074S, 1:2000).

## **Plasmid construction**

Constitutive active MEK (S218D, S222D) was constructed by site-directed mutagenesis. Evolved LOV (eLOV) was amplified from a synthetic gBlock from IDT based on a sequence from the previous work<sup>24</sup>. The sequence of LEXY was amplified from NLS-mCherry-LEXY (Addgene, catalog #72655). The full sequence of GLIMPSe was shown in Figure S8.

## **Cell culture and transfection**

HEK293T cells were cultured in DMEM supplemented with 10% fetal bovine serum and penicillinstreptomycin. PC12 cells were cultured in F12K medium supplemented with 15% horse serum and 2.5% FBS. All cell cultures were maintained in a standard incubator at 37 °C with 5% CO<sub>2</sub>. HEK293T and PC12 cells were seeded in 12 well plates, and cells were transfected once they reached 60% - 80% confluency. Transfection was performed using Turbofect transfection reagent following the vendor's instruction. For light-induced stabilization of MKP3 experiments, 100 ng CA MEK-EGFP, 200 ng EGFP-p2A-MKP3-eLOVtevS-deg and 700 ng of NLS-mCherry-TEV-LEXY plasmid were cotransfected in PC12 cells. For light-induced CA MEK stabilization experiments 50 ng of EGFP-p2A-CA MEK-eLOVtevS-deg and 950 ng of NLS-mCherry-TEV-LEXY plasmid was cotransfected in PC12 cells. After 3 hours of transfection, transfection medium was replaced with growth medium (F12K + 15% horse serum +2.5% FBS). After overnight recovery, the cell culture was exchanged to a low-serum medium (1.5% horse serum + 0.025% FBS) for another 24 hours to reduce the base-level ERK activity.

## Western Blot

After transfection, recovery, and starvation, cells were harvested and lysed with a mixture of RIPA buffer and protease/phosphatase inhibitor cocktail. Lysates were centrifuged, and supernatants were mixed with NuPAGE<sup>™</sup> LDS Sample Buffer and β-Mercaptoethanol. Samples were subjected to SDS-PAGE on a precast gel, followed by overnight transfer to PVDF membrane. Two different blots were run using the same cell lysate to probe the phospho-ERK and pan ERK, respectively.

#### Construction of a programmable LED device

For both long-term and short-term light illumination, the LED array was constructed by assembling a 6-by-4 blue LED array with 24 blue LEDs (B4304H96, Linrose Electronics) on a breadboard. LED intensity can be continuously tuned through a tunable voltage and a current-limiting resistor.

The breadboard was hosted in an aluminum box, and a light diffuser film was positioned above the LED array to make the light intensity homogeneous in the defined area. The light intensity at the cell culture plate was measured by a power meter (PM100D, S121C, Thorlabs).

#### **Statistical analysis**

The p-values were determined by performing two-tailed, unpaired t-test, or one-way ANOVA test using the GraphPad Prism software.



**Figure S1. Degron functions normally with the fusion protein HA-MKP3.** In PC12 cells, the fusion of degrons at the C-terminus of HA-MKP3 resulted in a 15.7-fold reduction of protein level compared to HA-MKP3 assayed by Western blot analysis with anti-HA antibody. Values were presented by mean  $\pm$  SD (n=2).



Figure S2. Absence of LEXY shows increased stabilization of MKP3 and reduced PC12 differentiation ratio in the dark. Differentiation ratio calculated for PC12 cells transfected with CA MEK-EGFP, EGFP-p2A-MKP3-eLOVtevS-3X degron, and TEV-mRuby2. Cells were illuminated with 0.5 mW/cm<sup>2</sup> blue light or kept in the dark for 45 h before imaging. Values represent the mean  $\pm$  SD of two biological replicates (n = 2) with more than 150 cells counted per replicate. Differentiation ratio= (# of transfected + differentiated cells) / # of transfected cells.



Figure S3. Absence of eLOV in MKP3-tevS-deg abolishes light-dark contrast for MKP3reduced PC12 cell differentiation. Differentiation ratio calculated for PC12 cells transfected with CA MEK-EGFP, mRuby2-p2A-MKP3-tevS-3X degron, and NLS-mCherry-TEV-LEXY. Cells were illuminated with 0.5 mW/cm<sup>2</sup> blue light or kept in the dark for 45 h before imaging. Values represent the mean  $\pm$  SD of two biological replicates (n = 2) with more than 150 cells counted per replicate. Differentiation ratio= (# of transfected + differentiated cells) / # of transfected cells.



**Figure S4.** The total amount of proteins remains consistent between the light-stimulated sample and the dark control upon inhibition of new protein synthesis with cycloheximide. PC12 cells transfected with CA MEK-EGFP, EGFP-p2A-HA-MKP3-eLOVtevS-3X degron, and NLS-mCherry-TEV-LEXY were treated with 100 ng/µL cycloheximide, a translation inhibitor, 24 h after transfection. Cell were incubated in cycloheximide for 18 h followed by light exposure for different time interval. The total amount of HA-tagged protein remained consistent across different conditions as new protein synthesis was blocked.



Figure S5. Light illumination for 30 min in PC12 cells transfected with GLIMPSe-CA MEK is sufficient to induce significant neurite outgrowth. CA-MEK rescued by 30 min light treatment shows a differentiation ratio of 0.6 compared with 0.2 for the dark control. The bar graph is presented with mean  $\pm$  SD averaged over three biological replicates (n = 3) with more than 200 cells counted per replicate.



Figure S6. Absence of eLOV in CA MEK-tevS-deg abolishes light-dark contrast for PC12 cell differentiation. Differentiation ratio calculated for PC12 cells transfected with mRuby2-p2A-CA MEK-tevS-3X degron and NLS-mCherry-TEV-LEXY. Cells were illuminated with 0.5 mW/cm<sup>2</sup> blue light or kept in the dark for 45 h before imaging. Values represent the mean  $\pm$  SD of three biological replicates (n = 3) with more than100 cells counted per replicate. Differentiation ratio= (# of transfected + differentiated cells) / # of transfected cells



**Figure S7. Long-term light exposure does not cause cell death.** (a) PC12 cells were exposed under blue light for 45-h ( $0.5 \text{ mW/cm}^2$ ) followed by co-staining with Calcein AM (staining live cells). Less than 1% dead cells were observed in both light and dark conditions. Values represent the mean ± SD of three biological replicates (n = 3) with more than 500 cells counted per replicate. (b) Representative fluorescent and phase-contrast images of PC12 cells in the same field of view under light and dark treatment. Scale bar: 50 µm.

1	TCT	'AGG	GCT	ACT	ACA	CTT	GAA	CGT	'ATT	'GAG	AAG	AGT	TTT	GTC	ATT	ACT	GAC	CCA	AGA	TTG
1	S	R	A	Т	Т	L	Е	R	Ι	Е	K	S	F	V	Ι	Т	D	Ρ	R	L
61	CCA	GAT	AAT	CCC	ATT	ATA	TTC	GTT	TCC	GAT	AGT	TTC	TTG	CAG	TTG	ACA	GAA	TAT	AGC	CGT
21	Ρ	D	Ν	Ρ	Ι	Ι	F	V	S	D	S	F	L	Q	L	Т	Ε	Y	S	R
121	GAA	GAA	ATT	TTG	GGA	AGA	AAC	TGC	AGG	TTT	CTA	CAA	GGT	CCT	GAA	ACT	GAT	CGC	GCG	ACA
41	Ε	Е	Ι	L	G	R	Ν	С	R	F	L	Q	G	Ρ	E	Т	D	R	A	Т
181	GTG	AGA	AAA	ATT	AGA	GAT	GCC	АТА	GAT	'AAC	CAA	ACA	GAG	GTC	ACT	GTT	CAG	CTG	ATT	AAT
61	V	R	K	Ι	R	D	A	Ι	D	Ν	Q	Т	Ε	V	Т	V	Q	L	Ι	Ν
241	TAT	ACA	AAG	AGT	GGT	AAA	AAG	TTC	TGG	AAC	CTC	TTT	CAC	TTG	CAG	CCT	ATG	CGA	GAT	CAG
81	Y	Т	K	S	G	K	K	F	W	Ν	L	F	Η	L	Q	Ρ	Μ	R	D	Q
301	AAG	GGA	GAT	GTC	CAG	TAC	TTT	ATT	'GGG	GTT	CAG	TTG	GAT	GGA	ACT	GAG	AGG	GTC	CGA	GAT
101	K	G	D	V	Q	Y	F	Ι	G	V	Q	L	D	G	Т	Е	R	V	R	D
361	GCT	GCC	GAG	AGA	GAG	GCT	GTC	ATG	CTG	GTT	AAG	AAA	ACT	GCA	GAA	GAA	ATT	GAT	GAG	GCG
121	А	A	Ε	R	Ε	A	V	М	L	V	K	K	Т	A	E	Е	Ι	D	Е	A
421	GCA	AAA	gaq	aac	ctq	tac	ttc	caq	ato	GGT	GGA	GGC	TCT	GGT	AGA	CTC	TAT	GAA	TTT	AGG
141	A	K	Е	Ν	L	Y	F	Q	М	G	G	G	S	G	R	L	Y	Ε	F	R
481	TTG	ATG	ATG	ACC	TTC	TCC	GGG	CTC	AAT	'CGC	GGT	TTT	GCA	TAC	GCA	.CGG	TAC	AGT	GGA	TCC
161	L	Μ	Μ	Т	F	S	G	L	Ν	R	G	F	A	Y	A	R	Y	S	G	S
541	GCT	AGC	GGT	AGA	CTC	TAT	GAG	TTT	'AGA	CTG	ATG	ATG	ACA	TTC	TCT	GGA	CTT	AAC	AGA	GGG
181	A	S	G	R	L	Y	Е	F	R	L	Μ	Μ	Т	F	S	G	L	Ν	R	G
601	TTC	GCC	TAT	GCC	CGA	TAT	TCT	GGA	TCC	GGT	AGG	CTT	TAT	GAG	TTT	CGC	CTG	ATG	ATG	ACA
201	F	A	Y	A	R	Y	S	G	S	G	R	L	Y	Ε	F	R	L	Μ	Μ	Т
661	TTT	TCC	GGG	TTG	AAC	AGG	GGC	TTC	GCT	'TAT	GCT	CGC	TAC	TCA	tag					
221	F	S	G	L	N	R	G	F	A	Y	A	R	Y	S	*					
	Evolved LOV (eLOV) TEV recognition site (tevS)																			
	Degron								Degron							Degron				

**Figure S8. DNA and amino acid sequences of GLIMPSe.** The domain of evolved LOV domain, tevS, and three codon-optimized degron are marked in distinct colors. The protein of target (e.g., MKP3 and CA MEK) were fused at the N-terminus of GLIMPSe.