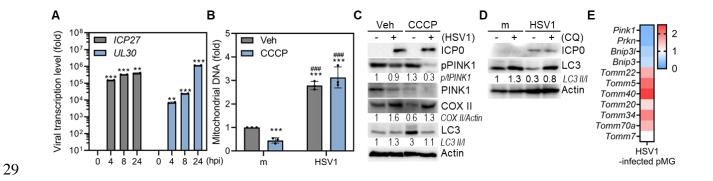
1	Supplementary materials
2	Pharmacological targeting of mitophagy via ALT001 improves herpes simplex
3	virus 1 (HSV1)-mediated microglial inflammation and promotes amyloid $\beta$
4	phagocytosis by restricting HSV1 infection
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8	Cho <sup>5</sup> , Ji-Yeun Hur <sup>6</sup> , Yongjin Yoo <sup>1, 3</sup> , Kihoon Han <sup>1, 3</sup> , Hosun Park <sup>4</sup> , Jeanho Yun <sup>2</sup> *, Ok Sarah
9	Shin <sup>1,7</sup> *
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25 26	* Co-correspondence: oshin@korea.ac.kr (OSS); Tel.: 82-2-2626-3280, yunj@dau.ac.kr (JY); Tel.:82-51-240-2919

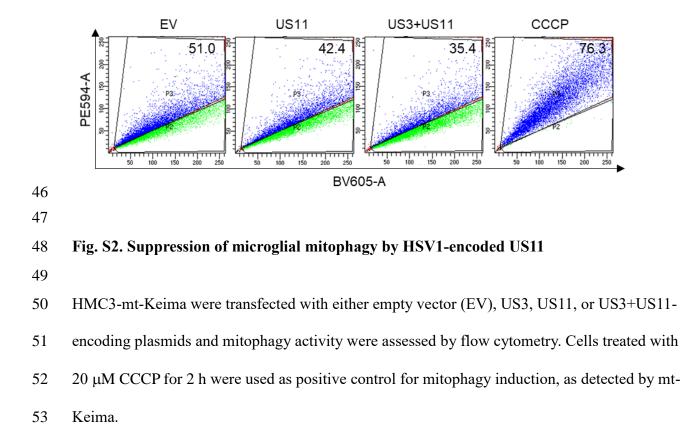
#### 28 Figure S1.

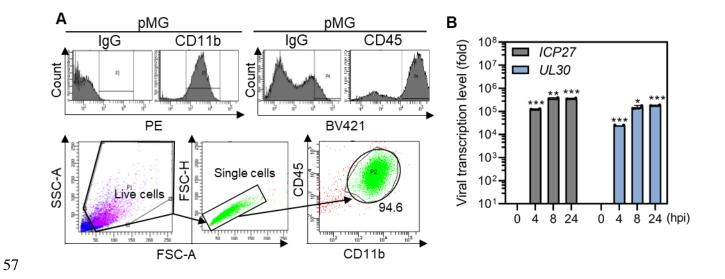


30 Fig. S1. Suppression of mitophagy by HSV1 in murine microglia

A Murine microglial cell line (BV2) were infected with HSV1 (MOI 1). At indicated time points, 31 32 RNA was isolated from cells, and HSV1 ICP27 and UL30 expression levels were analyzed by qRT-PCR. **B**, **C** BV2 were infected with HSV1 at MOI 1 for 24 h, followed by vehicle control 33 (Veh) or 20 µM CCCP treatment for 2 h. Mitochondrial DNA copies were quantified by PCR (B) 34 35 and an immunoblot was performed to determine protein expression of HSV1 ICP0, phospho-PINK1 (p-PINK1), PINK1, Mitochondria-encoded cytochrome C oxidase II (COX II), LC3, and 36 37 Actin (C). Band quantification relative to Actin is shown below the blot. **D** BV2 were infected 38 with HSV1 (MOI 1) for 24 h, followed by 10 µM chloroquine (CQ) treatment for 8 h. 39 Immunoblot analysis was performed to examine the levels of HSV1 ICP0, LC3, and Actin. E A 40 heatmap was generated using gene expression profiles from bulk RNA-sequencing of HSV1-41 infected pMG compared with mock-infected pMG (MOI 10). Data are presented as means  $\pm$  SD (n =3 biological replicates; p values were calculated by Student's t test). \*\* p < 0.01; \*\*\*p < 0.01; 42 0.001, compared with mock (m)-infected Veh-treated group.  $^{\#\#\#}p < 0.001$ , compared with mock-43 infected CCCP-treated group. 44

## **Figure S2.**



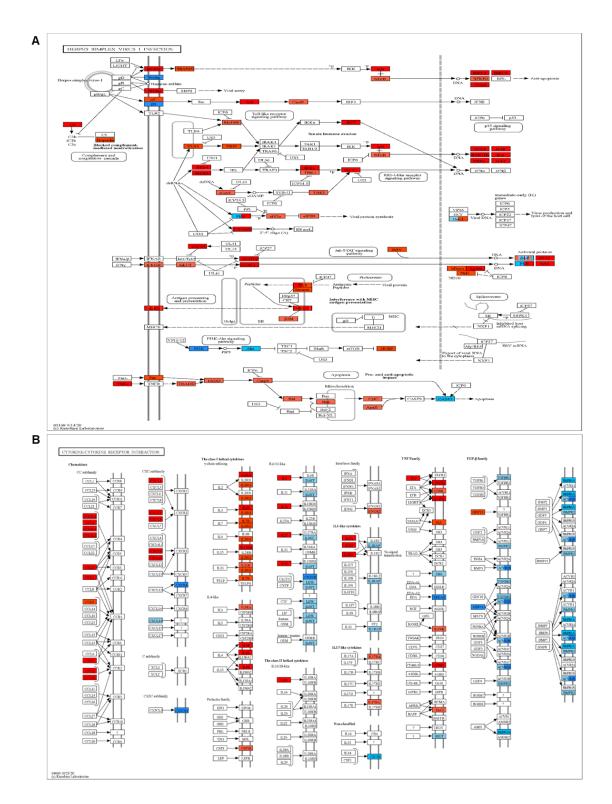


58 Fig. S3. Characterization of isolated primary murine microglia (pMG)

A Mixed glial cells (MxG) were isolated from mouse pups and grown for 14 days. After 14 days, primary murine microglia were isolated from MxG and stained with anti-CD11b and anti-CD45 antibodies for flow cytometry. **B** pMG were infected with HSV1 at MOI 10. At indicated time points, RNA was isolated from cells, and qRT-PCR was performed to examine the level of *HSV1 ICP27* and *UL30*. Data are presented as means  $\pm$  SD (n =3 biological replicates; *p* values were calculated by Student's t test). \*\* *p* < 0.01; \*\*\**p* < 0.001, versus mock (*m*)-infected group.

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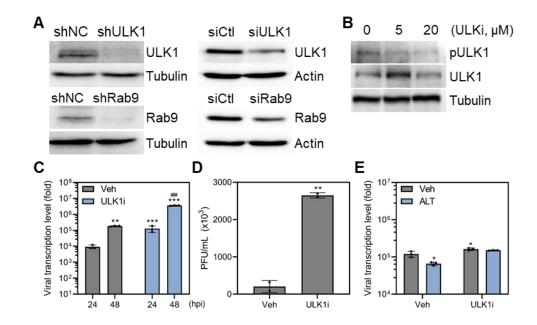
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# 70 Fig. S4. Alterations in HSV-1-associated host antiviral signaling pathway in microglia

71	A, B KEGG pathway map show upregulated (red) and downregulated (blue) genes in HSV1-
72	infected pMG (MOI 10) compared with mock-infected pMG. Herpes simplex virus 1 infection
73	(KEGG #; mmu05168) (A) and Cytokine-Cytokine receptor interaction (KEGG #; 04060) (B)
74	were generated.
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87 Figure S5.

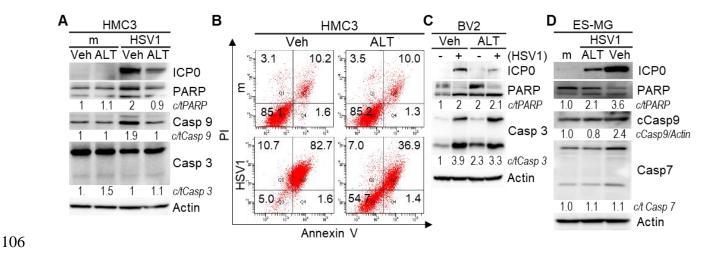






90 A HMC3 were transfected with scrambled negative control (shNC), ULK1-specific shRNA 91 (shULK1), and Rab9-specific shRNA (shRab9) or control siRNA (siCtl), ULK1-specific siRNA 92 (siULK1), and Rab9-specific siRNA (siRab9). The knockdown efficiency of shRNA or siRNA 93 was validated by immunoblot analysis showing protein levels of ULK1, Rab9, Actin, and Tubulin. B HMC3 were treated with 20 µM ULK1 inhibitor (SBI-0206965, ULK1i) for 48 h and 94 95 a representative immunoblot analysis of phospho-ULK1 (p-ULK1), ULK1, and Tubulin is shown. C, D HSV1 was inoculated in HMC3 (MOI 1) and cells were treated with 20 µM ULK1i. 96 97 HSV1 ICP27 mRNA levels were analyzed by qRT-PCR (C) and a plaque assay were performed to determine HSV1 titer (D). p < 0.05; p < 0.01; p < 0.01; p < 0.001, versus vehicle (Veh)-treated 98 group at 24hpi.  $^{\#}p < 0.01$ , versus Veh-treated at 48hpi. E HSV1 was inoculated in HMC3 at 99 100 MOI 1 and cells were treated with 30 µM ALT001 (ALT) and 20 µM ULK1i. At 48 hpi, RNA 101 was isolated from cells, and HSV1 ICP27 mRNA levels were analyzed by qRT-PCR. \*p < 0.05, 102 *versus Veh-treated group.* Data are presented as means  $\pm$  SD (n =3 biological replicates; p values

103 were calculated by Student's t test).

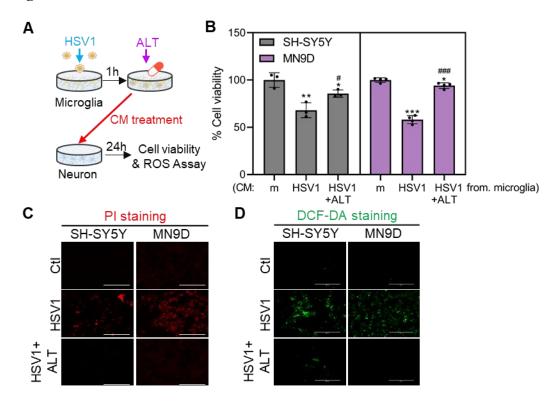




108 A, B HSV1 were inoculated in HMC3 (MOI 1) and cells were treated with vehicle (Veh) or 30 109 µM ALT001 (ALT). At 48 hpi, HSV1 ICP0, cleaved (c) PARP/total (t) PARP, cleaved caspase 110 9/total caspase 9 and cleaved caspase 3/total caspase 3 expression is shown by immunoblot 111 analysis (A). Actin was used as a protein loading control. Numbers below the blot represent 112 quantified band intensity by densitometric analysis. Cells were stained with Annexin V-FITC/PI 113 and apoptosis was analyzed by flow cytometry (B). C, D BV2 (C) or ES-MG (D) cells were 114 infected with HSV1 (MOI 1) in the presence or absence of 30 µM ALT. HSV1 ICP0, cleaved 115 PARP/total PARP, and cleaved caspase 3/total caspase 3 is shown by immunoblot analysis. 116 Images are representatives of three independent experiments. Band quantification relative to 117 Actin is shown below the blot.

119 Figure S7.

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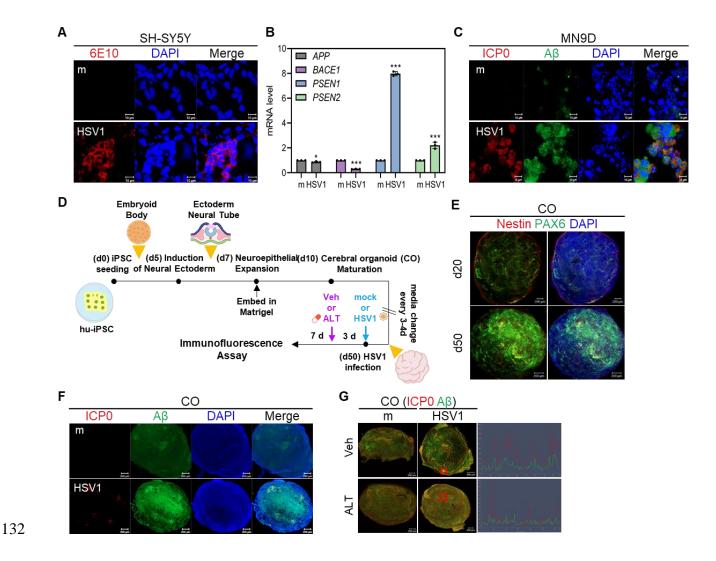


121 Fig. S7. ALT001 suppresses HSV1-mediated cell death in neuronal cells

122 A A schematic representation of conditioned media (CM) treatment from microglia to neuronal 123 cells is shown. **B** Human and mouse neuronal cells (SH-SY5Y and MN9D) were treated with 124 CM collected from mock (m)-, HSV1-, or HSV1-infected with ALT001 (HSV1+ALT) microglia. 125 After 24 h, CCK8 assay was performed to measure cell viability. Statistical analysis: one-way ANOVA with Dunnett's post-hoc correction. \*p < 0.05; \*\*p < 0.01; \*\*\*p < 0.001, versus CM 126 from m-infected group.  ${}^{\#}p < 0.05$ ,  ${}^{\#\#\#}p < 0.001$ , versus CM from Veh-treated HSV1-infected 127 128 group. C, D CM-treated neuronal cells were stained with propidium iodide (PI) (red) or 2',7'-129 Dichlorofluorescein Diacetate (DCF-DA) (green). Stained cells were analyzed by EVOS FL

130 Auto 2 Imaging system. Scale bar =  $200 \ \mu m$ .

### 131 **Figure S8.**



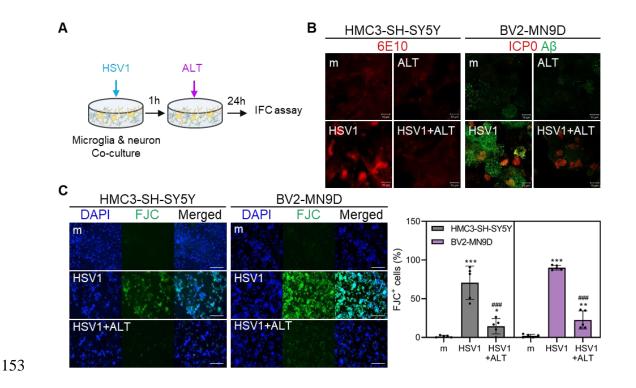


A, B Human neuroblastoma cells (SH-SY5Y) were infected with HSV1 (MOI 1) for 48 h and

135 stained against 6E10 (red) antibodies for confocal microscopy analysis (A) and mRNA levels of

- 136 amyloid beta precursor protein (APP), beta-secretase 1 (BACE1), presenilin 1 (PSEN1), and
- 137 *presenilin 2 (PSEN2)* was measured by qRT-PCR (means  $\pm$  SD; n = 3) (B). Statistical analysis;
- 138 Student's t test. \*p < 0.05; \*\*\*p < 0.001, versus mock (m)-infected group. Scale bar=10 µm. C

139	Mouse dopaminergic neuronal cells (MN9D) were infected with HSV1 (MOI 1) for 48 h and
140	were stained with HSV1 ICP0 (red) and A $\beta$ (green) for confocal microscopy analysis. Scale bar
141	=10 $\mu$ m. <b>D</b> Schematic representation of HSV1 infection in cerebral organoid (CO) model is
142	shown. E On day 20 or 50 after CO generation, CO were stained with neuronal marker, Nestin
143	(red) and PAX6 (green). Scale bar=200 $\mu$ m. F On day 50 of CO generation, CO were infected
144	with HSV1 (10 <sup>5</sup> PFU) for 3 days followed by immunofluorescence staining against HSV1 ICP0
145	(red) and A $\beta$ (green) for confocal microscopy analysis. Scale bar =200 $\mu$ m. G CO were infected
146	with HSV1 (10 <sup>5</sup> PFU) for 3 days and incubated with vehicle (Veh) or 30 $\mu$ M ALT001 (ALT) for
147	7 days. CO were stained with HSV1 ICP0 (red) or A $\beta$ (green) for confocal microscopy analysis.
148	The graph on the right represents the quantification of fluorescence intensity in the red box area
149	of images. Scale bar =200 $\mu$ m.



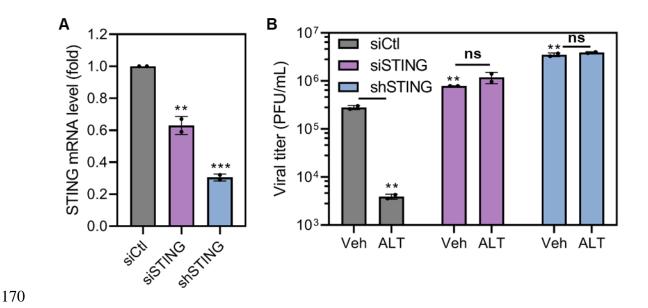
### 154 Fig. S9. ALT001 reduces degeneration of neurons in microglia-neuron co-culture model

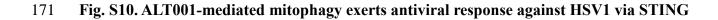
155 A A schematic representation of microglia & neuron co-culture system is shown. HSV1 was 156 inoculated in human microglia-neuron (HMC3-SH-SY5Y) or murine microglia-neuron (BV2-157 MN9D) co-culture model and these cells were treated with 30 µM ALT001 (ALT). B At 24 hpi, 158 cells were stained with 6E10 (red), A $\beta$  (green) or HSV1 ICP0 (red) for confocal microscopy 159 analysis. Scale bar =10  $\mu$ m. C cells were stained with FJC (green). % FJC-positive cells were 160 calculated using at least 5 different images. Scale bar=200 µm. Data are expressed as the means 161 ± SD of at least three independent experiments. Statistical analysis: one-way ANOVA with Dunnett's post-hoc correction. \*p < 0.05; \*\*p < 0.01; \*\*\*p < 0.001, versus vehicle (Veh)-treated 162 mock (m)-infected group.  ${}^{\#}p < 0.05$ ;  ${}^{\#\#\#}p < 0.001$ , versus Veh-treated HSV1-infected group. 163 164 165

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168 **Figure S10**.

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172 A STING mRNA expression level was measured in control siRNA-transfected HMC3 (siCtl),

173 STING-specific siRNA-transfected HMC3 (siSTING), and shSTING-stably expressing HMC3

174 (shSTING) by qRT-PCR. B STING knockdown cells by siRNA or shRNA introduction were

175 infected with HSV1 at MOI 1 in the presence or absence of vehicle (Veh) or 30 µM ALT001

176 (ALT). Supernatant was collected at 48 hpi and a plaque assay was performed in Vero cells using

177 the supernatant. Statistical analysis: one-way ANOVA with Dunnett's post-hoc correction.

178 \*\**p*<0.01; \*\*\**p* < 0.001, versus siCtl-transfected Veh group. ns=non-significant.