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OPEN AMPK contributes to autophagosome maturation and lysosomal fusion

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AMP-activated protein kinase (AMPK) regulates autophagy initiation when intracellular ATP level decreases. However, the role of AMPK during autophagosome maturation is not fully understood. Here, we report that AMPK contributes to efficient autophagosome maturation and lysosomal fusion. Using CRISPR-Cas9 gene editing, we generated AMPK α 1 knockout HEK293T cell lines, in which starvationinduced autophagy is impaired. Compound C, an AMPK-independent autophagy inducer, and trehalose, an mTOR-independent autophagy inducer were used to examine the role of AMPK in autophagosome maturation and lysosomal fusion. While the treatment of control cells with either compound C or trehalose induces activation of autophagosomes as well as autolysosomes, the treatment of AMPK α 1 knockout cells with compound C or trehalose induces mainly activation of autophagosomes, but not autolysosomes. We demonstrate that this effect is due to interference with the fusion of autophagosomes with lysosomes in AMPK $\alpha 1$ knockout cells. The transient expression of AMPK $\alpha 1$ can rescue autophagosome maturation. These results indicate that AMPK $\alpha 1$ is required for efficient autophagosome maturation and lysosomal fusion.

Autophagic flux is the entire process of macroautophagy (hereafter referred to as autophagy), ranging from the inclusion of cargo within the autophagosome to digestion in the autolysosome, and either increased autophagic flux or a block in autophagic flux can result in autophagosome accumulation. During the process of increased autophagic flux, the autophagosome fuses with the lysosome to form an autolysosome, which provides an acidic environment for lysosomal hydrolases to destroy the cargo molecules^{2,3}. Autophagosome maturation and the lysosomal fusion process can be analyzed by tandem fluorescent-tagged LC3 (ptf-LC3) or the level of p62/ SQSTM1^{2,4,5}

AMP activated protein kinase (AMPK) is a crucial cellular energy sensor protein and is activated by a low energy state in the cell^{6,7}. The AMPK complex consists of catalytic α subunits and regulatory β and γ subunits, and the mammalian genome has multiple AMPK subunit isoforms $(\alpha 1, \alpha 2, \beta 1, \beta 2, \gamma 1, \gamma 2, \gamma 3)^8$. The expression of AMPK $\alpha 1$ complex is ubiquitous; however, the expression of AMPK $\alpha 2$ is high in skeletal muscle, the heart, and the liver 9,10. AMPK is one of the major autophagy regulators, and the role of AMPK in autophagy initiation has been clearly shown. Under glucose starvation, AMPK associates with and activates autophagy-initiating kinase Ulk1, which is an orthologue of yeast ATG1, the most upstream component of the autophagy machinery¹¹⁻¹³. In addition, the activation of AMPK can phosphorylate TSC2 and the activated TSC2 can suppress mTOR complex 1 (mTORC1) to induce autophagy 14,15. However, the role of AMPK in autophagosome maturation and lysosome fusion is not fully understood. Several reports have suggested that AMPK is involved in autophagosome maturation. Although AMPK can negatively regulate mTORC1 signaling and mTORC1 activation can suppress autophagosome maturation via UVRAG phosphorylation 16,17, the relationship between AMPK and activation of autophagosome maturation is not clear. Metformin, an activator of AMPK, can induce autophagy, as can compound C, an inhibitor of AMPK¹⁸⁻²⁰. Compound C induced autophagosome formation in an AMPK-independent manner, since neither the AMPK activator, AICAR nor metformin blocked compound C-induced autophagosome formation¹⁹.

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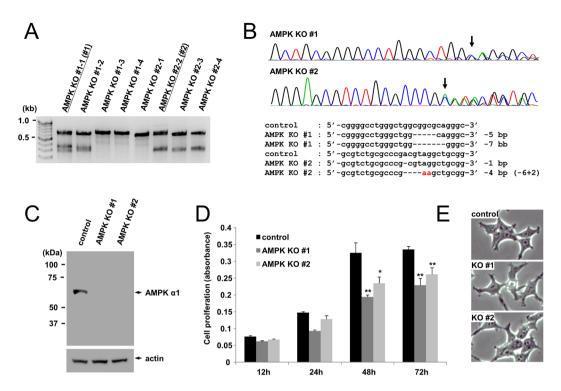


Figure 1. Generation of AMPK α 1 knockout (KO) HEK293T cells. (**A**) Validation of AMPK α 1 KO by T7 endonuclease 1 (T7E1) assay. HEK293T cells were transfected with either pX459/AMPK α 1 gRNA #1 or pX459/AMPK α 1 gRNA #2, and single colonies were isolated. The genomic PCR products were analyzed by T7E1 assay. Two different clones (KO #1 and KO #2) were used for the further experiments. (**B**) DNA sequencing analysis revealed the presence of AMPK α 1 mutation in KO #1 and KO #2 (Top). The CRISPR-Cas9 system introduced an indel mutation in the target sites of the AMPK α 1 gene (bottom). (**C**) Validation of AMPK α 1 KO by Western blotting. Equal amounts of HEK293T wild-type (WT) and AMPK α 1 KO cell lysates were subjected to Western blotting with an anti-AMPK α 1 antibody. Experiments were repeated three times with similar results. (**D**) Proliferation of AMPK α 1 KO cell lines was analyzed by MTT assay. The MTT assay was performed in triplicate, and the graph shows the average and standard deviation (SD). Control vs. knockout cells. *P < 0.05, **P < 0.01. (**E**) Phase contrast microscope images (400x) of control cells and AMPK α 1 knockout cells.

Trehalose, a disaccharide present in non-mammalian species, inhibits solute carrier $2\,A$ (SLC2A) and induces an mTOR independent autophagy $^{21-23}$.

In this report, we generated AMPK $\alpha 1$ knockout cell lines, which impaired starvation-induced autophagy. Because the transfection efficiency of HEK293T cells is high, knockout HEK293T cells were used for transient expression experiments involving the autophagy marker and cell signaling reporter. Compound C and trehalose treatment induced autophagosome formation in both control and AMPK $\alpha 1$ knockout cells. However, autophagosome maturation and lysosome fusion were blocked in AMPK $\alpha 1$ knockout cells. The overexpression of AMPK rescued AMPK function, indicating that AMPK is required for efficient autophagic flux even though compound C-induced autophagosome formation is AMPK independent.

Results

Generation of AMPK α 1 knockout (KO) HEK293T cells. We generated AMPK α 1 knockout (KO) cell lines using the CRISPR-Cas9 gene editing system²⁴. Two AMPK α 1 guide RNA sets were synthesized and cloned into a pX459 vector. AMPK α 1 knockout plasmids were transfected into HEK293T cells. After selection, we isolated single colonies and analyzed the insertion or deletion mutation (indel) using T7 endonuclease 1 (T7E1) assays (Fig. 1A). Next, we analyzed the indel mutation of the PCR products of target DNA by nucleotide sequencing and confirmed that the AMPK α 1 gene was mutated (Fig. 1B). Finally, we demonstrated that the expression of AMPK α 1 protein was abolished in HEK293T cells by Western blotting (Fig. 1C). These results collectively indicate that AMPK α 1 knockout cell lines were successfully established by the CRISPR-Cas9 system. Because gene knockout often affects cell proliferation, we examined the cell proliferation of AMPK α 1 knockout cells by MTT assay. Although there was no remarkable phenotypic change, the proliferation of AMPK α 1 knockout cells was significantly reduced by up to 25% compared to HEK293T control cells (Fig. 1D,E).

AMPK α **1** knockout impairs starvation-induced autophagy. Many reports support that AMPK is required for autophagy initiation upon cellular starvation 12,25 . We examined whether AMPK α 1 is required for autophagy using AMPK α 1 knockout cells. To examine the regulation of autophagy by AMPK, we used an mRFP-GFP-LC3 reporter construct (ptfLC3), which appears yellow when autophagosomes form and red when autolysosomes form². Control cells and AMPK α 1 knockout cells were incubated with HBSS for 1 h and

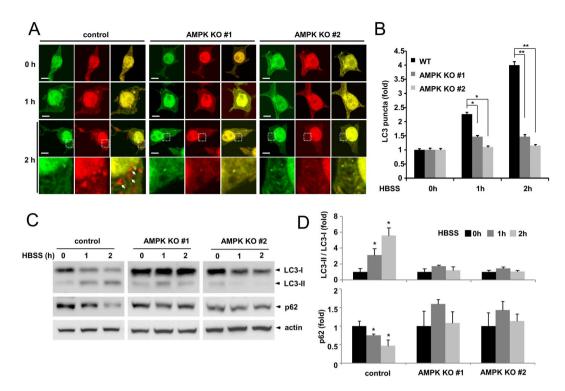


Figure 2. AMPK $\alpha 1$ knockout inhibits autophagy initiation upon starvation. (A) Control cells and AMPK $\alpha 1$ knockout cells were transfected with a plasmid encoding mRFP-GFP-LC3, and cells were either mock-treated or starved in HBSS medium for the indicated time (1 h or 2 h). Bars: $10\,\mu m$. (B) The numbers of LC3 puncta in control cells and AMPK $\alpha 1$ knockout cells were counted (n = 20). Control cells vs. AMPK knockout cells, *P < 0.05, **P < 0.005. (C) AMPK $\alpha 1$ knockout represses starvation-induced autophagy. Control cells and AMPK $\alpha 1$ knockout cells were incubated in HBSS medium, and the cell lysates were probed with anti-LC3 and anti-p62 antibodies. (D) The LC3-II, LC3-I and p62 bands were quantified, and the relative expression levels of LC3-II/LC3-I and p62 are shown in the graph. 0 h vs. 1 h or 2 h, *P < 0.05.

2 h, respectively, and cells were observed under a fluorescence microscope. HEK293T control cells showed an increased number of yellow and red puncta (autophagosome/autolysosome), However, AMPK α 1 knockout cells showed only a limited number of yellow puncta, indicating that autophagy initiation was impaired (Fig. 2A,B). We also examined the autophagy markers LC3 and p62. In control cells, the level of LC-II/LC3-I protein was gradually elevated, and the level of p62 was reduced, however, AMPK α 1 knockout cells did not show any significant changes (Fig. 2C,D). These results confirm that AMPK α 1 is required for efficient autophagy initiation in HEK293T cells.

Repression of autophagic flux in AMPK \alpha 1 knockout cells. A previous study demonstrated that compound C, an inhibitor of AMPK, can induce autophagy in an AMPK-independent manner¹⁹. For this reason, we examined whether compound C could induce autophagy in AMPK al knockout cells. Control cells and AMPK α1 knockout cells were transfected with an mRFP-GFP-LC3 reporter construct (ptfLC3), and cells were treated with compound C for 18 h. Both control cells and AMPK α1 knockout cells showed cytoplasmic puncta, suggesting that autophagosomes had formed. While control cells showed red puncta (autolysosomes), AMPK α1 knockout cells showed yellow puncta (autophagosomes), indicating that autophagic flux was impaired in AMPK α1 knockout cells (Fig. 3A,B). We also examined the expression levels of autophagy markers LC3 and p62. While the level of p62 protein in HEK293T control cells was decreased with compound C treatment, this decrease was less prominent in AMPK α1 knockout cells (Fig. 3C,D). In addition, the treatment of a lysosomal inhibitor, bafilomycin A1 in combination with compound C showed that AMPK α 1 knockout cells showed a decreased difference in p62 level between bafilomycin A1 treatment and untreated samples. This indicates a decreased autophagic flux in AMPK α1 knockout cells (Figs S1 and S2). Overall, these results indicate that autophagic flux in AMPK al knockout cells was impaired with compound C treatment. We also examined the level of LC3-II protein. Both HEK293T control cells and AMPK α1 knockout cells showed elevated levels of LC3-II, confirming that autophagosomes were formed in both cell lines (Fig. 3C). However, the level of LC3-II was not significantly changed in AMPK α 1 knockout cells (Fig. S3). These results collectively indicate that AMPK α 1 is not required for autophagosome formation after treatment with compound C, but is necessary for the successful formation of autolysosomes that occurs as a result of autophagic flux.

Disruption of AMPK α **1 interferes with the formation of autolysosomes.** To form mature autolysosomes, autophagosomes need to be fused with lysosomes. Because AMPK α 1 knockout cells did not form autolysosomes with compound C treatment (Fig. 3A), we examined whether the autophagosomes could fuse with

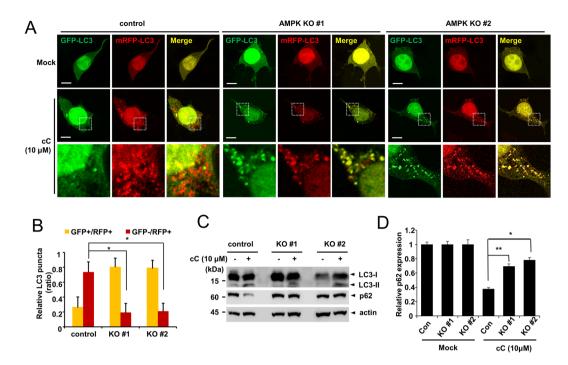


Figure 3. AMPK $\alpha 1$ knockout represses autophagic flux. (A) Control cells and AMPK $\alpha 1$ knockout cells were transfected with a plasmid encoding mRFP-GFP-LC3, and cells were either mock-treated (vehicle alone) or treated with compound C (cC, $10\,\mu\text{M}$) for $18\,\text{h}$. Bars: $10\,\mu\text{m}$. (B) Ratio quantification of autophagosomal LC3 puncta to autolysosomal LC3 puncta (n = 10). Control cells vs. knockout cells, *P < 0.0001. (C,D) HEK293T cells and AMPK $\alpha 1$ knockout cells were treated with compound C ($10\,\mu\text{M}$) for $18\,\text{h}$, and the cell lysates were probed with anti-LC3 and anti-p62 antibodies. The p62 bands were quantified, and the relative expression levels are shown in the graph. Control cells vs. AMPK $\alpha 1$ knockout cells, *P < 0.05, **P < 0.005.

lysosomes in AMPK $\alpha 1$ knockout cells. Since GFP-LC3 protein loses its fluorescence in low lysosomal pH, we used mRFP-LC3 for the detection of autophagosomes and autolysosomes. Control cells and AMPK $\alpha 1$ knockout cells were transfected with mRFP-LC3. Twenty-four h after transfection, the cells were treated with compound C, followed by immunostaining with anti-LAMP1 antibody (a lysosomal marker). We found many mRFP-LC3 puncta co-stained with anti-LAMP1 in control cells; however, mRFP-LC3 puncta were positive for LAMP1 staining in only a few AMPK $\alpha 1$ knockout cells (Fig. 4A,B). These results suggest that the formation of mature autolysosomes was impaired in AMPK $\alpha 1$ knockout cells due to disruption of the fusion of autophagosomes with lysosomes. In order to confirm that AMPK $\alpha 1$ knockout interferes with autolysosome formation, we employed transmission electron microscopy to assess the ultra-structure of autophagosomes and autolysosomes. Control cells and AMPK $\alpha 1$ knockout cells were treated with compound C for 18 h, and then cells were fixed and examined with a transmission electron microscope. While the control cells showed autolysosomes as well as autophagosomes, the AMPK $\alpha 1$ knockout cells showed mainly autophagosomes and lysosomes (Fig. 4C,D). These results collectively indicate that AMPK is involved in autophagosome maturation and lysosome fusion.

Disruption of AMPK α **1 inhibits AMPK-related signaling.** We examined AMPK activity in AMPK α 1 knockout cells that were treated with compound C, a selective AMPK inhibitor. AMPK is the main regulator of acetyl-CoA carboxylase (ACC), which can be phosphorylated by AMPK activation²⁶. Control cells and AMPK α 1 knockout cells were either mock treated or treated with compound C, and the cell lysates were probed with an anti-phospho-ACC antibody (p-ACC). While the level of phospho-ACC was decreased in control cells with compound C treatment, phospho-ACC was hardly detectable in AMPK α 1 knockout cells (Fig. 5A). These results demonstrate that AMPK α 1 is required for the phosphorylation of ACC. In addition, we examined cell viability upon treatment with compound C. Cell viability was decreased by compound C in a dose-dependent manner, However, the cell viability of AMPK knockout cells was not clearly different from that of the control cells (Fig. 5B,C).

Transient expression of AMPK α 1 rescues the AMPK α 1 knockout phenotype. Because AMPK α 1 knockout cells showed impairment in autophagosome maturation, we examined whether the overexpression of AMPK α 1 can rescue AMPK activity and autophagosome maturation. AMPK α 1 knockout cells were transfected with an AMPK α 1 plasmid encoding full-length AMPK α 1 cDNA and an mRFP-GFP-LC3 plasmid, and the cells were then treated with compound C. Transient expression of AMPK α 1 resulted in the presence of autolysosome puncta (red), indicating efficient autophagic flux (Fig. 6A,B). In addition, the restoration of AMPK α 1 resulted in the phosphorylation of ACC (Fig. 6C). Finally, expression of AMPK α 1 resulted in a reduction in

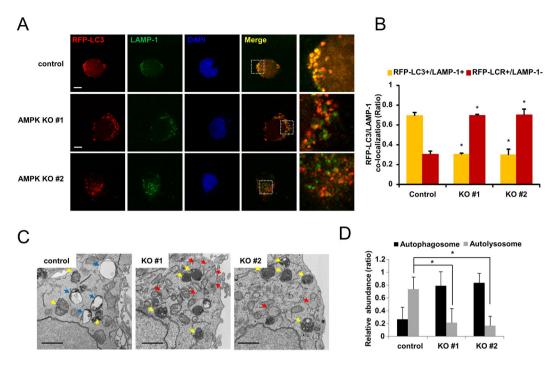


Figure 4. AMPK α 1 knockout interferes with the formation of autolysosomes. (A) Control cells and AMPK α 1 knockout cells were transfected with the plasmid encoding mRFP-LC3 and treated with compound C for 18 h. Cells were fixed and stained with an anti-LAMP1 antibody. Bars: $10\,\mu\text{m}$. (B) Ratio quantification of autophagosomal LC3 puncta (RFP-LC3 only) to autolysosomal LC3 puncta (RFP-LC3 and LAMP-1) Each data represents the mean and standard deviation of two independent experiments (n = 30), control cells vs AMPK Knockout cells, *P<0.05. (C) Representative electron micrograph of HEK293T cells and AMPK α 1 knockout cells after compound C treatment. Yellow arrows denote the autophagosome; blue arrows, autolysosomes; and red arrows, lysosomes. Bars: $1\,\mu\text{m}$. (D) Ratio quantification of autophagosomal puncta to autolysosomal puncta. We counted autophagosomes and autolysosomes in 54 individual cells (n = 54) of each group, and calculated the relative abundance. Graph shows the average and standard deviations. Control cells vs. knockout cells, *P<0.0001.

p62 level after compound C treatment (Fig. 6C). These results collectively indicate that AMPK α 1 expression can rescue the AMPK α 1 knockout phenotype.

AMPK $\alpha 1$ knockout represses trehalose induced autophagic flux. Since the AMPK inhibitor compound C induces autophagy independent of AMPK, yet regulates autophagic flux AMPK-dependently, we sought to clarify the role of AMPK in autophagic flux by inducing autophagy in another manner. A previous study showed that trehalose can induce autophagy in mTOR independent manner²¹. Control cells and AMPK $\alpha 1$ knockout cells were transfected with an mRFP-GFP-LC3 reporter construct (ptfLC3), and cells were treated with trehalose. Both control cells and AMPK $\alpha 1$ knockout cells showed cytoplasmic puncta, suggesting that autophagosomes had formed. While control cells showed mainly red puncta (autolysosomes), AMPK $\alpha 1$ knockout cells showed yellow puncta (autophagosomes) (Fig. 7A). We also examined the expression levels of autophagy markers LC3 and p62. While the level of p62 protein in control cells was decreased with trehalose treatment, the level of p62 protein was significantly increased in AMPK $\alpha 1$ knockout cells (Fig. 7B,C). The level of LC3-II was not significantly changed in AMPK $\alpha 1$ knockout cells (Fig. S4).

To further confirm these results, we examined whether autophagosomes could fuse with lysosomes in AMPK α 1 knockout cells upon trehalose treatment. Control cells and AMPK α 1 knockout cells were transfected with mRFP-LC3. Twenty-four hours after transfection, the cells were treated with trehalose, followed by immunostaining with anti-LAMP1 antibody. We found that autolysosome formation is impaired in AMPK α 1 knockout cells (Fig. 7D,E). Finally, we performed a rescue experiment by expressing AMPK α 1 in AMPK α 1 knockout cells. The overexpression of AMPK α 1 restores the autophagosome maturation by forming autolysosomes (Fig. 8A,B). We also examined p62 levels and show that the restoration of AMPK α 1 decreased the level of p62 upon trehalose treatment (Fig. 8C). These results indicate that AMPK α 1 knockout does not disrupt the trehalose-induced formation of autophagosomes, but does delay autolysosome formation by inhibiting autophagosome maturation.

Discussion

Here, we report that AMPK increases autophagic flux by contributing to autophagosome maturation and autolysosome fusion. The role of AMPK in autophagy initiation has been clearly shown in previous studies ^{12,13}; however, the role of AMPK in the later stages of autophagy, autophagosome maturation, and autolysosome fusion has not been characterized. We showed that AMPK $\alpha 1$ depletion impaired the progression of autophagy induced

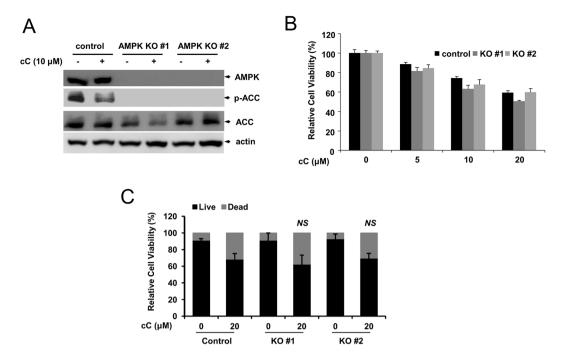


Figure 5. AMPK $\alpha 1$ knockout deregulates AMPK-related signaling. (A) AMPK $\alpha 1$ knockout decreases the phosphorylation level of acetyl-CoA carboxylase (ACC). Control cells and AMPK $\alpha 1$ knockout cells were either mock-treated (vehicle alone) or treated with compound C (cC, $10\,\mu\text{M}$) for $18\,\text{h}$, and the equal amount of cell lysates were probed with the indicated antibodies. (B) Proliferation of AMPK $\alpha 1$ KO cell lines after compound C treatment. HEK293T control and knockout clones were seeded into 24-well plates and incubated with the indicated concentrations of compound C. MTT assays were performed in triplicate, and the graph shows the average and standard deviation (SD). (C) Cell viability of AMPK $\alpha 1$ KO cell lines were measured with Trypan blue assay after compound C treatment. Control cells vs. knockout cells, NS: not significant.

by compound C treatment. This indicates that AMPK $\alpha 1$ contributes to the progression of autophagy, as well as autophagy initiation.

We generated AMPK α 1 knockout cell lines using the CRISPR-Cas9 gene editing method in HEK293T cells. Although there are two isoforms of the AMPK α subunit, AMPK α 1 and AMPK α 2, the knockout of only AMPK α 1 impaired autophagosome formation under starvation in HEK293T cells, suggesting that AMPK α 1 plays an important role in starvation-induced autophagy in HEK293T cells (Fig. 2). Since we generated the knockout cells with the HEK293T cell line, in which the transfection efficiency is high, we obtained definitive results using the autophagy marker mRFP-GFP-LC3. In addition, we used two different guide RNAs to avoid biased results due to off-target effects.

It is controversial whether compound C promotes or inhibits autophagy. It has been reported that compound C inhibited autophagy by suppressing AMPK activity in U2OS osteosarcoma cells, while another study reported that compound C induced autophagy in U251 glioma cells by an AMPK-independent pathway^{19,27}. In a previous report that claimed that compound C induced autophagy by an AMPK-independent pathway, the authors did not examine the later stages of autophagy, such as autophagosome maturation¹⁹. Here, we demonstrated that compound C induced autophagosome formation via an AMPK-independent manner. However, cells treated with compound C require AMPK for the maturation of autophagosomes and for autolysosome formation. Therefore, our experimental results provide another explanation for compound C-induced autophagy.

Since compound C inhibits AMPK activity, it is quite paradoxical that AMPK is involved in compound C-induced autophagy by promoting autophagosome maturation. We used a high concentration of compound C (10 μ M) and found that the AMPK activity is significantly reduced. Therefore, our results imply that this reduced AMPK activity is sufficient for autophagosome maturation and autolysosome formation.

Because compound C is also an AMPK inhibitor, it is difficult to make a clear conclusion about the role of AMPK $\alpha 1$ in autophagy from these experiments. However, we also showed that autophagosome formation was normal and autolysosome formation was impaired in AMPK $\alpha 1$ knockout cells in the presence of trehalose, an mTOR-independent autophagy inducer. Taking both the trehalose and compound C experiments together, we conclude that AMPK $\alpha 1$ is required for efficient autophagosome and autolysosome formation during autophagy.

Although autophagosome maturation and autolysosome formation are significantly impaired in AMPK α 1 knockout cells, we did not find a significant change in cell viability upon treatment with compound C (Fig. 5B,C). These results are consistent with previously published results that apoptosis induced by compound C is AMPK independent^{28,29}. In this respect, autophagy induced by compound C is neither cytoprotective nor cytotoxic, but rather nonprotective^{4,30}. Compound C has been reported to sensitize cancer cells to drug-mediated apoptosis^{31,32}. Therefore, compound C might sensitize tumor cells to other cancer drugs by regulating autophagy.

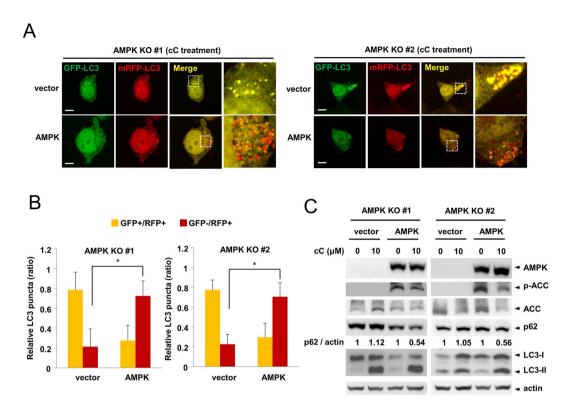


Figure 6. Expression of AMPK α 1 recues the AMPK α 1 knockout phenotype. (A) AMPK α 1 knockout cells were transfected with mRFP-GRP-LC3 in the presence or absence of the AMPK plasmid. Twenty-four h after transfection, cells were treated with compound C for 18 h. Bars: $10\,\mu\text{m}$. (B) Ratio quantification of autophagosomal LC3 puncta to autolysosomal LC3 puncta (n = 10). Vector vs AMPK α 1 plasmid, *P < 0.0001. (C) AMPK α 1 knockout cells were transfected with the plasmid encoding AMPK α 1. Twenty-four hour after transfection, cells were treated with compound C for 18 h. Equal amounts of cell lysates were probed with the indicated antibodies.

In this report, we showed that AMPK expression is required for efficient autophagosome maturation and lysosome fusion in AMPK $\alpha 1$ knockout cells. Since we tested our hypothesis in specific conditions and in a specific cell line, the requirement of AMPK for efficient autophagosome maturation and lysosome fusion might not apply to other conditions. Therefore, further experiments will be required to confirm our results in other cell lines and other conditions.

Materials and Methods

Cell culture and cell proliferation assay. HEK293T cells were grown in DMEM medium (Welgene, Korea) supplemented with 10% fetal bovine serum (Gibco, Waltham, MA, USA) and 1% antibiotic-antimycotic solution (Welgene, Seoul, Korea). Cell proliferation was measured using the [4,5-dimethylthiazol-2-yl]-2,5-diphenyltrazolium bromide (MTT) assay. Briefly, cells were seeded in a 24-well plate and then incubated overnight. At the indicated time, MTT solution was added to a final concentration of 1 mg/ml and incubated for 3 additional hours. Trypan blue assay was performed to measure cell viability. Cells were stained with trypan blue solution and the numbers of live cells and dead cells were counted by a TC10 automated cell counter (Bio-Rad, Richmond, CA, USA). MTT was purchased from USB Corporation (Cleveland, OH, USA) and Compound C from Sigma (St. Louis, MO, USA). Compound C was dissolved in DMSO (stock solution, 10 mM) and control cells were treated with an equal volume of DMSO (vehicle alone).

Generation of the AMPK α **1 knockout cell line with CRISPR/Cas9.** Guide RNA sequences for use in the CRISPR/Cas9 system were designed at the CRISPR design website (http://crispr.mit.edu/), provided by the Feng Zhang Lab²⁴. The sequences for the insert oligonucleotides for human AMPK α 1 gRNA #1 and #2 were as follows 5'-CACCGA AGATCGGCCACTACATTC-3'/5'-AAACGAATGTAGTGGCCGATCTTC-3'and 5'-CACCGCCGAGAAGCAGAAACACGA-3'/5'-AAACTCGTGTTTCTGGTTCTCGGC-3', respectively. The AMPK α 1 guide RNA targets exon 1 of the AMPK α 1 gene. The complementary oligonucleotides for the guide RNAs (gRNAs) were cloned into a pX459 CRISPR/Cas9-Puro vector (Addgene, Cambridge, MA, USA). HEK293T cells were transfected with lipofectamine transfection reagent (Invitrogen, Carlsbad, CA, USA). Two days after transfection, cells were treated with 1 μg/ml of puromycin (Sigma) for three days. After two weeks, colonies were isolated, and the AMPK α 1 sequences were amplified by PCR and analyzed using the T7 endonuclease (T7E1) assay, DNA sequencing (Macrogen, Seoul, Korea), and Western blotting. The T7E1 enzyme was purchased from New England Biolabs (Beverly, MA, USA).

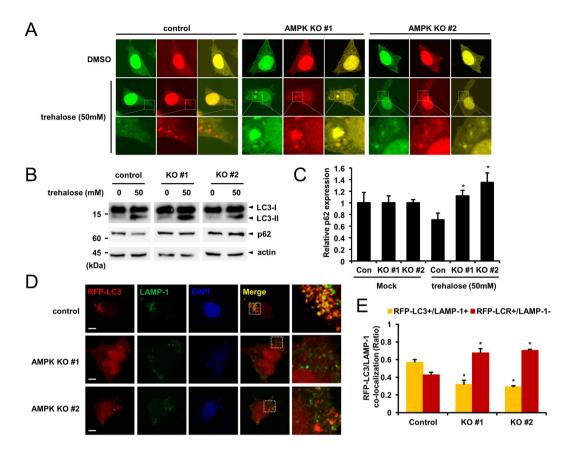


Figure 7. AMPK α 1 knockout represses trehalose induced autophagic flux. (A) Control cells and AMPK α 1 knockout cells were transfected with mRFP-GRP-LC3. Twenty-four h after transfection, cells were either mock-treated (vehicle only) or treated with trehalose (50 mM) for 4 h. (B,C) HEK293T cells and AMPK α 1 knockout cells were treated with trehalose (50 mM) for 6 h, and the cell lysates were probed with anti-LC3 and anti-p62 antibodies. The p62 bands were quantified, and the relative expression levels are shown in the graph. Control cells vs. AMPK α 1 knockout cells, *P < 0.05. (D,E) Cells were transfected with mRFP-LC3 plasmid. 24 h after, cells were treated with Trehalose for 6 h. Cells were fixed and stained with anti-LAMP1 antibody. Ratio quantification of autophagosomal LC3 puncta (RFP-LC3 only) to autolysosomal LC3 puncta (RFP-LC3 and LAMP-1). Each data represents the mean and standard deviation of two independent experiments (n = 30), control cells vs AMPK Knockout cells, *P < 0.05.

Western blotting. For Western blot analysis, polypeptides in whole cell lysates were resolved by SDS-PAGE and transferred to nitrocellulose membrane filters. Proteins were detected with a 1:1000 or 1:5000 dilution of primary antibody using an enhanced chemiluminescence (ECL) system. Images were acquired using the Chemidoc-it 410 imaging system (UVP, Upland, CA, USA) and LAS4000 system (GE Healthcare, Uppsala, Sweden). The following primary antibodies were used: anti-AMPK α 1 (Cell Signaling Technology, Beverly, MA, USA), anti-phospho-AMPK α 1 (Cell Signaling Technology), anti-LC3 (MBL international, Watertown, MA, USA), anti-p62 (MBL international), anti-acetyl-CoA carboxylase (ACC) (Cell Signaling Technology), anti-phospho acetyl-CoA carboxylase (p-ACC) (Cell Signaling Technology), and anti-actin (ABM, Richmond, BC, Canada).

Immunofluorescence and confocal microscopy. HEK293T wild-type and AMPK knockout cells were grown on sterilized glass coverslips. After plasmid transfection, cells were fixed with 4% paraformaldehyde. For immunostaining, cells were blocked with 10% goat serum (Gibco) in phosphate-buffered saline (PBS; Wellgene, Seoul, Korea), stained with a 1:500 dilution of primary antibody in PBS, and then stained with a 1:1000 dilution of fluorescence-conjugated secondary antibody (Invitrogen, Carlsbad, MA, USA). Finally, slides were washed three times with PBS and mounted in mounting medium containing DAPI (Vector Laboratories, Burlingame, CA, USA). Images were captured with a Carl Zeiss LSM710 confocal microscope (Oberkochem, Germany). The plasmid pTF-LC3 was purchased from Addgene (Cambridge, MA, USA) and the anti-LAMP1 antibody from Santa Cruz Biotechnology (Santa Cruz, CA, USA).

Transmission electron microscopy. HEK293T cells were sequentially fixed with 2.5% glutaraldehyde and 1% osmium tetroxide on ice for 2 hours and washed with PBS. The tissues were then serially dehydrated in ethanol and propylene oxide, embedded in an Epon 812 mixture, and polymerized in an oven at 70 °C for 24 hours. The sections acquired from the polymerized blocks were collected on grids, counterstained with uranyl acetate

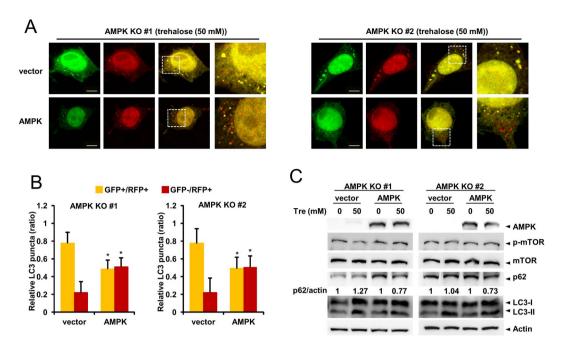


Figure 8. Expression of AMPK α 1 recovers trehalose induced autophagic flux. (A,B) AMPK α 1 knockout cells were transfected with mRFP-GFP-LC3 in the presence or absence of the AMPK plasmid. Twenty-four hour after, cells were treated with trehalose for 6 h. Ratio quantification of autophagosomal LC3 (GFP+/RFP+) puncta to autolysosomal LC3 puncta (GFP-/RFP+) Bars: $10\,\mu\text{m}$. (n = 15, vector vs AMPK plasmid, *P<0.0001). (C) AMPK α 1 knockout cells were transfected with AMPK plasmid. Twenty-four hour after transfection, cells were treated with trehalose for 6 h. Cell lysates were probed with the indicated antibody.

and lead citrate, and examined with a Bio-HVEM system (JEM-1400Plus at 120 kV and JEM-1000BEF at 1000 kV, JEOL, Tokyo, Japan).

Autophagosome was identified by its contents (morphologically intact organelles) with the existence of double membranes (partly visible), and autolysosome was identified by its contents (partially degraded cellular organelles due to cellular autolysis) with the clear empty space. Lysosome can be identified by its homogenous content.

Statistical methods. The results of the western blot, TEM data and LC3 puncta analysis were evaluated by a 2-tailed t test using Excel software (Microsoft, Seattle, WA, USA). P < 0.05 was considered significant.

Data availability. The datasets generated during and/or analyzed during the current study are available from the corresponding author on reasonable request.

References

- 1. Klionsky, D. J. et al. Guidelines for the use and interpretation of assays for monitoring autophagy (3rd edition). Autophagy 12, 1–222, https://doi.org/10.1080/15548627.2015.1100356 (2016).
- 2. Kimura, S., Noda, T. & Yoshimori, T. Dissection of the autophagosome maturation process by a novel reporter protein, tandem fluorescent-tagged LC3. *Autophagy* 3, 452–460, https://doi.org/10.4161/auto.4451 (2007).
- 3. Mizushima, N. & Yoshimori, T. How to interpret LC3 immunoblotting. *Autophagy* 3, 542–545, https://doi.org/10.4161/auto.4600 (2007).
- 4. Bjorkoy, G. et al. p62/SQSTM1 forms protein aggregates degraded by autophagy and has a protective effect on huntingtin-induced cell death. J Cell Biol 171, 603–614, https://doi.org/10.1083/jcb.200507002 (2005).
- 5. Namkoong, S. *et al.* The integral membrane protein ITM2A, a transcriptional target of PKA-CREB, regulates autophagic flux via interaction with the vacuolar ATPase. *Autophagy* 11, 756–768, https://doi.org/10.1080/15548627.2015.1034412 (2015).
- 6. Hardie, D. G., Ross, F. A. & Hawley, S. A. AMPK: a nutrient and energy sensor that maintains energy homeostasis. *Nat Rev Mol Cell Biol* 13, 251–262, https://doi.org/10.1038/nrm3311 (2012).
- Biol 15, 251–262, https://doi.org/10.1058/htm5511 (2012).

 Hardie, D. G. AMPK: a key regulator of energy balance in the single cell and the whole organism. Int J Obes (Lond) 32(Suppl 4), 87–12, https://doi.org/10.1038/iio.2008.116 (2008).
- 8. Hardie, D. G., Schaffer, B. E. & Brunet, A. AMPK: An Energy-Sensing Pathway with Multiple Inputs and Outputs. *Trends Cell Biol* 26, 190–201, https://doi.org/10.1016/j.tcb.2015.10.013 (2016).
- 9. Kemp, B. E. et al. Dealing with energy demand: the AMP-activated protein kinase. Trends Biochem Sci 24, 22–25, https://doi.org/10.1016/S0968-0004(98)01340-1 (1999).
- 10. Guo, Y. et al. AMP-activated kinase alpha2 deficiency protects mice from denervation-induced skeletal muscle atrophy. Arch Biochem Biophys 600, 56–60, https://doi.org/10.1016/j.abb.2016.04.015 (2016).
- Egan, D., Kim, J., Shaw, R. J. & Guan, K. L. The autophagy initiating kinase ULK1 is regulated via opposing phosphorylation by AMPK and mTOR. Autophagy 7, 643–644, https://doi.org/10.4161/auto.7.6.15123 (2011).
- 12. Kim, J., Kundu, M., Viollet, B. & Guan, K. L. AMPK and mTOR regulate autophagy through direct phosphorylation of Ulk1. *Nat Cell Biol* 13, 132–141, https://doi.org/10.1038/ncb2152 (2011).
- 13. Roach, P. J. AMPK > ULK1 -> autophagy. Mol Cell Biol 31, 3082–3084, https://doi.org/10.1128/MCB.05565-11 (2011).
- 14. Tripathi, D. N. et al. Reactive nitrogen species regulate autophagy through ATM-AMPK-TSC2-mediated suppression of mTORC1. Proc Natl Acad Sci USA 110, E2950–2957, https://doi.org/10.1073/pnas.1307736110 (2013).

- 15. van Veelen, W., Korsse, S. E., van de Laar, L. & Peppelenbosch, M. P. The long and winding road to rational treatment of cancer associated with LKB1/AMPK/TSC/mTORC1 signaling. *Oncogene* 30, 2289–2303, https://doi.org/10.1038/onc.2010.630 (2011).
- Dunlop, E. A. & Tee, A. R. The kinase triad, AMPK, mTORC1 and ULK1, maintains energy and nutrient homoeostasis. Biochem Soc Trans 41, 939–943, https://doi.org/10.1042/BST20130030 (2013).
- 17. Kim, Y. M. et al. mTORC1 phosphorylates UVRAG to negatively regulate autophagosome and endosome maturation. Mol Cell 57, 207–218, https://doi.org/10.1016/j.molcel.2014.11.013 (2015).
- 18. Shi, W. Y. et al. Therapeutic metformin/AMPK activation blocked lymphoma cell growth via inhibition of mTOR pathway and induction of autophagy. Cell Death Dis 3, e275, https://doi.org/10.1038/cddis.2012.13 (2012).
- Vucicevic, L. et al. Compound C induces protective autophagy in cancer cells through AMPK inhibition-independent blockade of Akt/mTOR pathway. Autophagy 7, 40–50, https://doi.org/10.4161/auto.7.1.13883 (2011).
- Tomic, T. et al. Metformin inhibits melanoma development through autophagy and apoptosis mechanisms. Cell Death Dis 2, e199, https://doi.org/10.1038/cddis.2011.86 (2011).
- Sarkar, S., Davies, J. E., Huang, Z., Tunnacliffe, A. & Rubinsztein, D. C. Trehalose, a novel mTOR-independent autophagy enhancer, accelerates the clearance of mutant huntingtin and alpha-synuclein. *J Biol Chem* 282, 5641–5652, https://doi.org/10.1074/jbc. M609532200 (2007).
- 22. Mayer, A. L. et al. SLC2A8 (GLUT8) is a mammalian trehalose transporter required for trehalose-induced autophagy. Sci Rep 6, 38586, https://doi.org/10.1038/srep38586 (2016).
- DeBosch, B. J. et al. Trehalose inhibits solute carrier 2A (SLC2A) proteins to induce autophagy and prevent hepatic steatosis. Sci Signal 9, ra21, https://doi.org/10.1126/scisignal.aac5472 (2016).
- Ran, F. A. et al. Genome engineering using the CRISPR-Cas9 system. Nat Protoc 8, 2281-2308, https://doi.org/10.1038/nprot.2013.143 (2013).
- 25. Rabinowitz, J. D. & White, E. Autophagy and metabolism. Science 330, 1344-1348, https://doi.org/10.1126/science.1193497 (2010).
- Sim, A. T. & Hardie, D. G. The low activity of acetyl-CoA carboxylase in basal and glucagon-stimulated hepatocytes is due to
 phosphorylation by the AMP-activated protein kinase and not cyclic AMP-dependent protein kinase. FEBS Lett 233, 294–298,
 https://doi.org/10.1016/0014-5793(88)80445-9 (1988).
- 27. Xu, Z. X. et al. A plant triterpenoid, avicin D, induces autophagy by activation of AMP-activated protein kinase. Cell Death Differ 14, 1948–1957, https://doi.org/10.1038/sj.cdd.4402207 (2007).
- 28. Liu, X., Chhipa, R. R., Nakano, I. & Dasgupta, B. The AMPK inhibitor compound C is a potent AMPK-independent antiglioma agent. *Mol Cancer Ther* 13, 596–605, https://doi.org/10.1158/1535-7163.MCT-13-0579 (2014).
- Jin, J. et al. AMPK inhibitor Compound C stimulates ceramide production and promotes Bax redistribution and apoptosis in MCF7 breast carcinoma cells. J Lipid Res 50, 2389–2397, https://doi.org/10.1194/jlr.M900119-JLR200 (2009).
- 30. Gewirtz, D. A. Cytoprotective and nonprotective autophagy in cancer therapy. *Autophagy* 9, 1263–1265, https://doi.org/10.4161/auto.25233 (2013).
- 31. Jang, J. H. et al. Compound C sensitizes Caki renal cancer cells to TRAIL-induced apoptosis through reactive oxygen species-mediated down-regulation of c-FLIPL and Mcl-1. Exp Cell Res 316, 2194–2203, https://doi.org/10.1016/j.yexcr.2010.04.028 (2010).
- 32. Kim, H. S. et al. Quercetin enhances hypoxia-mediated apoptosis via direct inhibition of AMPK activity in HCT116 colon cancer. Apoptosis 17, 938–949, https://doi.org/10.1007/s10495-012-0719-0 (2012).

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Author Contributions

M.J., R.P., J.P. designed the experiments. M.J., R.P., H.K., D.J., Y.H.H. performed the experiments. M.J., J.P. wrote the manuscript. S.N., I.S.J., J.I.L. reviewed and edited the manuscript.

Additional Information

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