

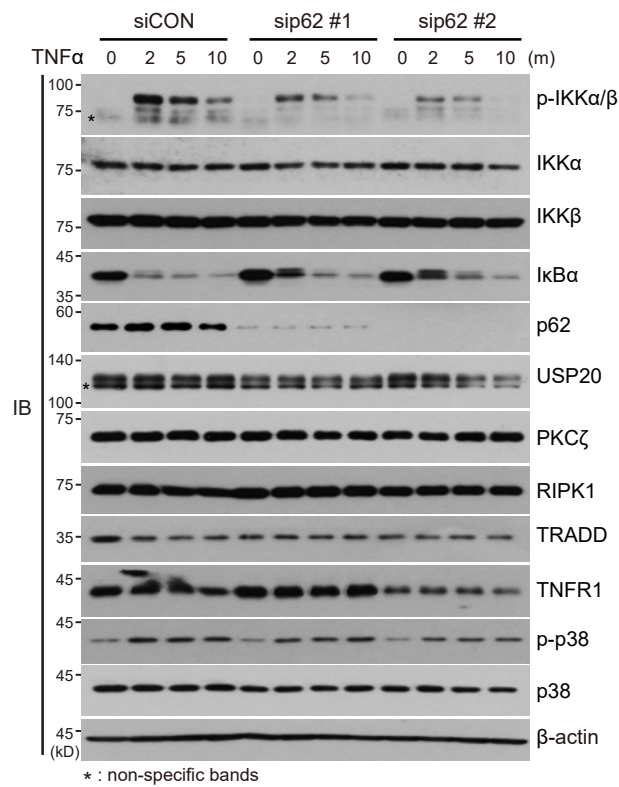
## **Supplementary Materials**

**The deubiquitinating enzyme USP20 regulates TNF $\alpha$ -induced NF- $\kappa$ B signaling pathway through the stabilization of p62**

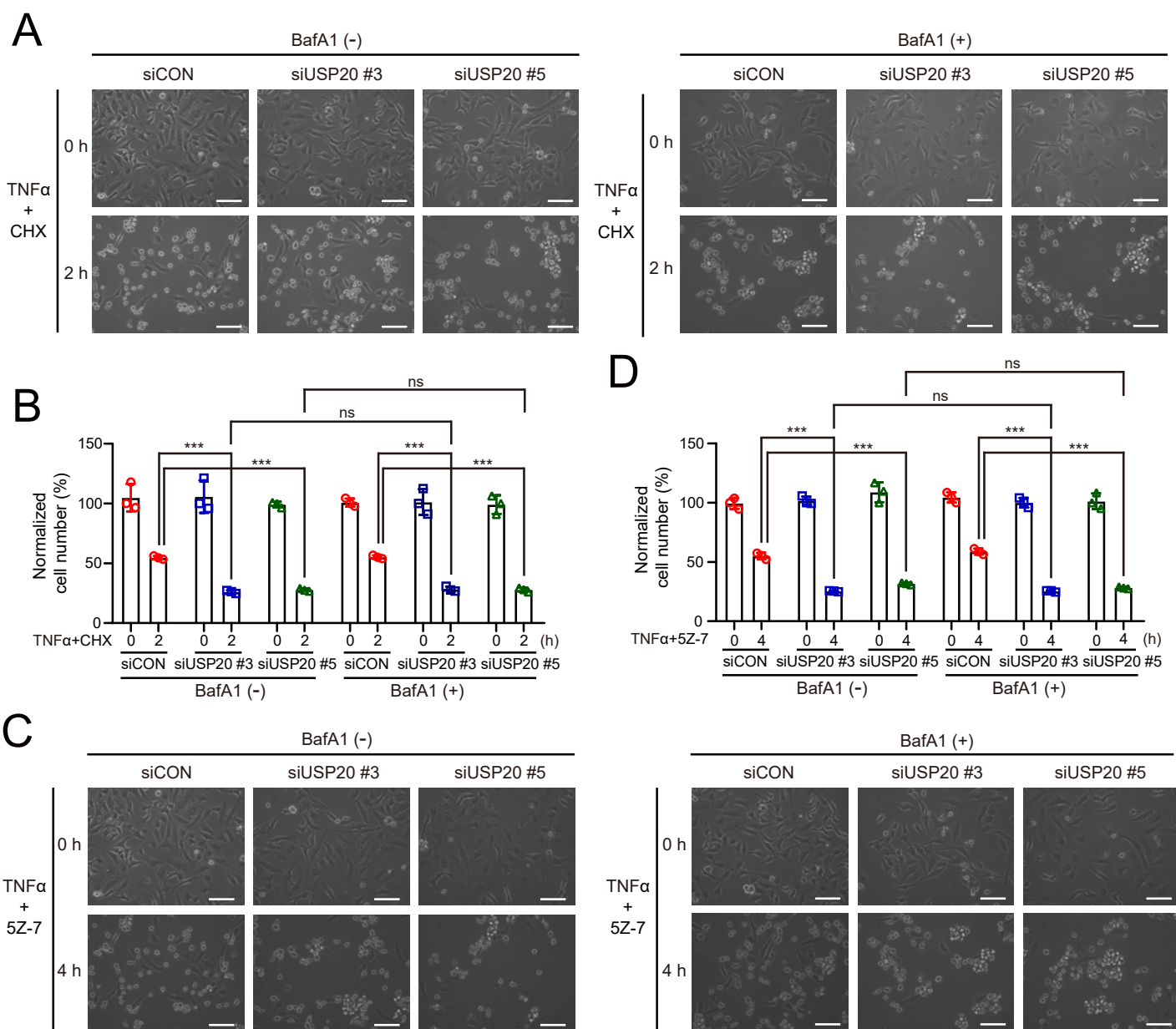
Jihoon Ha, Minbeom Kim, Dongyeob Seo, Jin Seok Park, Jaewon Lee, Jinjoo Lee, Seok Hee Park

### **<Table of Contents>**

1. Supplementary figures; Figure S1 – S3
2. Supplementary tables; Table S1 and S2

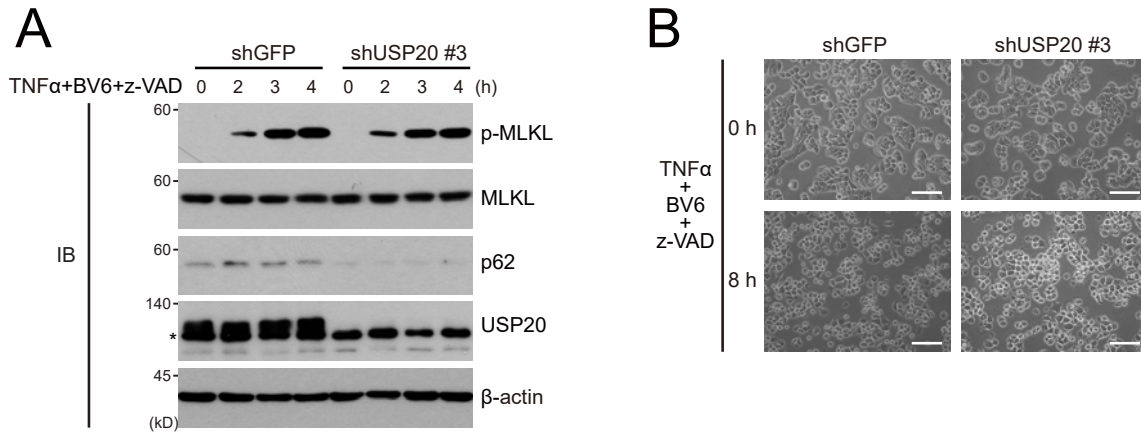


**Figure S1.** p62 depletion decreases TNF $\alpha$ -induced NF- $\kappa$ B signaling. HeLa cells were reverse-transfected with 20 nM control siRNA (siCON) or two independent p62-specific siRNAs (sip62 #1 and sip62 #2). Cells were treated with 10 ng/ml TNF $\alpha$  for the indicated time points. Total cell lysates were immunoblotted with the indicated antibodies. Expression of  $\beta$ -actin was used as a loading control. The immunoblot image is representative of three independent experiments.



**Figure S2.** Autophagy is not related to TNF $\alpha$ -mediated apoptosis under USP20 depletion.

**A, B:** USP20-depleted and control (siCON) HeLa cells were pre-treated with 100 nM bafilomycin A1 (BafA1) and subsequently treated with 20 ng/ml TNF $\alpha$  plus 10  $\mu$ g/ml cycloheximide (CHX) for 2 h. Morphological changes were observed by light microscopy (**A**). Scale bars, 1000  $\mu$ m. Live cell countings (**B**) were performed to measure cell viability. **C, D:** USP20-depleted and control (siCON) HeLa cells were pre-treated with 100 nM bafilomycin A1 (BafA1) and subsequently treated with 20 ng/ml TNF $\alpha$  plus 1  $\mu$ M 5Z-7 (TAK1 inhibitor) for 4 h. Morphological changes were observed by light microscopy (**C**). Scale bars, 1000  $\mu$ m. Live cell countings (**D**) were performed to measure cell viability. The data in this figure were statistically analyzed by two-way ANOVA followed by Bonferroni's multiple comparison test ( $***P < 0.001$  compared to the control cells without TNF $\alpha$ /CHX or TNF $\alpha$ /5Z-7, ns; not significant,  $n = 3$ ). The bars represent the mean  $\pm$  SD. The images in (**A**) and (**C**) are representative of three independent experiments.



**Figure S3.** USP20 depletion is not related to necroptosis.

**A, B:** USP20-depleted HT29 cells were generated by infections of lentiviruses expressing specific shRNA against endogenous *USP20* mRNA. As a negative control, lentiviruses expressing shRNA against *GFP* mRNA (shGFP) were used. USP20-depleted and control (shGFP) HT29 cells were treated together with 30 ng/ml TNF $\alpha$ , 1  $\mu$ M BV6 and 20  $\mu$ M z-VAD-fmk for 8 h. Cell lysates were immunoblotted with the indicated antibodies (**A**). Morphological changes were observed by light microscopy (**B**). Scale bars, 1000  $\mu$ m. The images in this figure are representative of three independent experiments.

**Table S1. Primer sequences used for real-time RT-PCR in this study.**

Gene	Species	Direction	Sequences
<i>GAPDH</i>	Human	Forward	5'-TGTAGTTGAGGTCAATGAAGGG-3'
		Reverse	5'-ACATCGCTCAGACACCATG-3'
<i>USP20</i>	Human	Forward	5'-CAATGGGCAGTGGTACGAGT-3'
		Reverse	5'-CCGCGAAGGTGTTGAACTTG-3'
<i>p62</i>	Human	Forward	5'-GTGGCTGTAACCTGCTGGAT-3'
		Reverse	5'-CTCTTTCAGGGACAGGCTGG-3'
<i>BFL1</i>	Human	Forward	5'-AGGTCCAAGCAAACGTCCA-3'
		Reverse	5'-ATCCACATCCGGGGCAATTT-3'
<i>cFLIP</i>	Human	Forward	5'-CATAAGCCGTTTGACCACGC-3'
		Reverse	5'-TGAACCGCTTCACGCCTAAT-3'

**Table S2. The sequences of siRNAs used in this study.**

<b>siRNA</b>	<b>Species</b>	<b>Sequences</b>
sip62 #1	Human	Sense : 5'- CAUGUCCUACGUGAAGGAUGAUU-3' Antisense : 5'-AAUCAUCCUUCACGUAGGACAUG-3'
sip62 #2	Human	Sense : 5'-GCAUUGAAGUUGAUAUUCGAUUU-3' Antisense : 5'-AAAUCGAUAUCAACUCAAUGC-3'
siUSP20 #3	Human	Sense : 5'-GCGAGUGGCUCAACAAGUU-3' Antisense : 5'-AACUUGUUGAGCCACUCGC-3'
siUSP20 #5	Human	Sense : 5'-GCCAGAACGUGAUCAAUGG-3' Antisense : 5'-CCAUUGAUCACGUUCUGGC-3'