

Supplementary Information

IKK ϵ isoform switching governs the immune response against EV71 infection

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Section details	Title	Page
Methods	Droplet digital PCR analysis	3
Tables		
Supplementary Table 1	Enriched pathways with isoform switching genes in EV71 infection	4
Supplementary Table 2	List of primers and probes	5
Supplementary Table 3	The raw data of ddPCR	6
Figures		
Supplementary Figure 1	The pathway enriched by isoform switching genes in EV71 infection	7
Supplementary Figure 2	IKK ϵ v2 transcriptionally induces ISGs expressions in EV71 infection through IRF7	10
Supplementary Figure 3	EV71 infection promotes ubiquitination	12
Supplementary Figure 4	IKK ϵ v2 increases IRF7 phosphorylation and IRF7 translocation in the presence of ubiquitin	14
Supplementary Figure 5	The interaction of IKK isoforms and IRF7	17
Supplementary Figure 6	The expression levels of HA-ubiquitin, referred to Figure 5	19
Supplementary Figure 7	Full-scan images of representative Western blots, referred to Figure 1	22
Supplementary Figure 8	Full-scan images of representative Western blots, referred to Figure 2	23
Supplementary Figure 9	Full-scan images of representative Western blots, referred to Figure 3	24
Supplementary Figure 10	Full-scan images of representative Western blots, referred to Figure 4	25
Supplementary Figure 11	Full-scan images of representative Western blots, referred to Figure 5	26
Supplementary Figure 12	Full-scan images of representative Western blots, referred to Supplementary Figure 2	28
Supplementary Figure 13	Full-scan images of representative Western blots, referred to Supplementary Figure 3	29
Supplementary Figure 14	Full-scan images of representative Western blots, referred to Supplementary Figure 4	30
Supplementary Figure 15	Full-scan images of representative Western blots, referred to Supplementary Figure 5	31
Supplementary Figure 16	Full-scan images of representative Western blots, referred to Supplementary Figure 6	32

Supplementary Methods

Droplet digital PCR analysis

Three commercially available probes purchased from Thermo Fisher were used to detect IKK ϵ isoforms in which Hs01063858-m1 detects all the three IKK ϵ isoforms, Hs01069870_m1 detects both IKK ϵ v1 and v2 and Hs01063855_g1 detects both IKK ϵ v1 and v3. After ddPCR assays, we subtracted copies of Hs01069870_m1 (detecting v1 and v2) from Hs01063858-m1 (detecting v1, v2, and v3) to have IKK ϵ v3. Similarly, we subtracted copies of Hs01063855_g1 (detecting v1 and v3) from Hs01063858-m1 (detecting v1, v2, and v3) to have IKK ϵ v2. Finally, copies of Hs01063858-m1 minus IKK ϵ v3 and IKK ϵ v2 leaves IKK ϵ v1. Ratio of individual IKK ϵ isoform was calculated by copy number of each IKK ϵ isoform divided by total copies of three isoforms.

Supplementary Table 1. Enriched pathways with isoform switching genes in EV71 infection

Rank	Pathway	Entities	Genes	<i>p</i> value
1	Toll-like receptor signaling pathway	6	IRAK4, AKT1, IKBKE, MYD88, IRF3, SPP1	0.022
2	PPAR signaling pathway	5	PPARD, DBI, SCP2, ACAA1, ANGPTL4	0.024
3	Neurotrophin signaling pathway	6	IRAK4, AKT1, BDNF, CAMK2B, SH2B1, TP73	0.047
4	Apoptosis	5	IRAK4, AKT1, TNFRSF10B, MYD88, IL1RAP	0.049

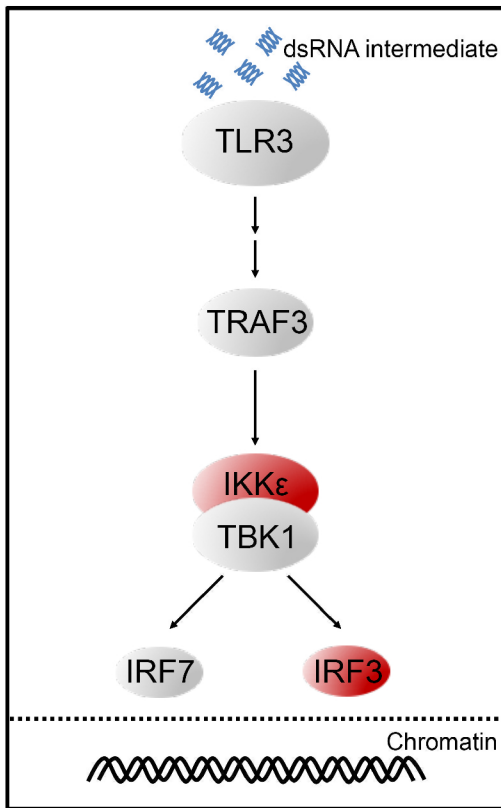
Supplementary Table 2. List of primers and probes

Gene name	Sequence (5'-3')
IFNb-SYBR-F	ATTGCCTCAAGGACAGGATG
IFNb-SYBR-R	GGCCTTCAGGTAATGCAGAA
OAS1-SYBR-F	CCCCATTATTGAAAAGTACCTGAGA
OAS1-SYBR-R	GCCGGGTCCAGGATCAC
ISG56-SYBR-F	CAGAACGGCTGCCTAATTTACA
ISG56-SYBR-R	GTGGGTCTGCTTTTTCTCTGT
ISG20-SYBR-F	CCCTGCGGGTGCTGAGT
ISG20-SYBR-R	TGTCCAAGCAGGCTGTTCTG
Mxa-SYBR-F	GCTACTGTGGCCAGAAAATC
Mxa-SYBR-R	TCATACTGGCTGCACAGGTTGT
Promoter Luc reporter	Sequence (5'-3')
OAS1-F	CCCGGTACCCTTAACAAAAAGAAAAGAGAC
OAS1-R	TTTAAGCTTTTTACCACCTTGGACACACA
ISG56-F	TAAGGTACCGCACCCAGCCAAGAATCATT
ISG56-R	CGCAAGCTTAGATCTGGCTATTCTGTCTT
ISG20-F	AAAGGTACCCCAAATCCCCTTGGTGAAA
ISG20-R	AAAAAGCTTCTCTCACCTGCCTGCCTCTG
Mxa-F	ACCGGTACCCCAAAGCTCACCAGTATCAA
Mxa-R	ATAAAGCTTCTCTGCTACCAGGCTGAGGA
Expression vector	Sequence (5'-3')
IRF7-HindIII-F	AAAAAGCTTATGGCCTTGGCTCCTGAGAGGGCAG
IRF7-BamHI-R	AAAGGATCCGGCGGGCTGCTCCAGCTCCATAAGG
Digital Probe	Target exon boundary
Hs01063858_m1	NM_014002.3 Exon 21-22
Hs01069870_m1	NM_014002.3 Exon 2-3
Hs01063855_g1	NM_014002.3 Exon 19-20

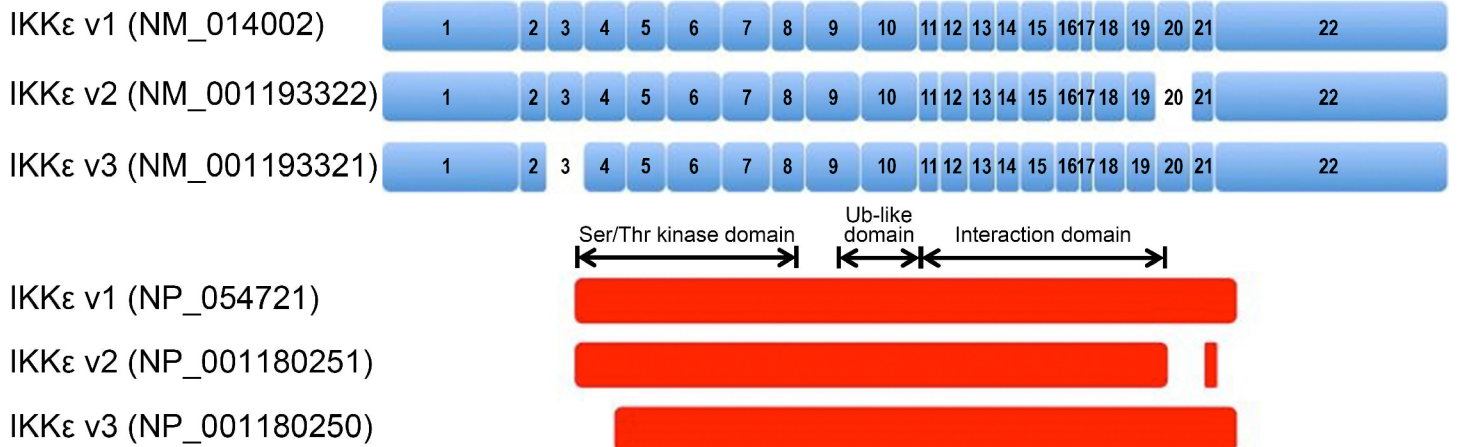
Supplementary Table 3. The raw data of ddPCR

Sample	Probes		
	Hs01063858	Hs01069870	Hs01063855
RD_Mock infection	3330	2960	2780
RD_4 h.p.i.	5140	4450	3820
RD_8 h.p.i.	1750	1436	1026
SH-SY5Y_Mock infection	2620	2120	2440
SH-SY5Y_12 h.p.i.	3260	2120	2900
SH-SY5Y_24 h.p.i.	2440	1656	2140
Copies of each IKKϵ			
	IKK ϵ v1	IKK ϵ v2	IKK ϵ v3
RD_Mock infection	2410	550	370
RD_4 h.p.i.	3130	1320	690
RD_8 h.p.i.	712	724	314
SH-SY5Y_Mock infection	1940	180	500
SH-SY5Y_12 h.p.i.	1760	360	1140
SH-SY5Y_24 h.p.i.	1356	300	784
Ratio of each IKKϵ			
	IKK ϵ v1	IKK ϵ v2	IKK ϵ v3
RD_Mock infection	72%	17%	11%
RD_4 h.p.i.	61%	26%	13%
RD_8 h.p.i.	41%	41%	18%
SH-SY5Y_Mock infection	74%	7%	19%
SH-SY5Y_12 h.p.i.	54%	11%	35%
SH-SY5Y_24 h.p.i.	56%	12%	32%

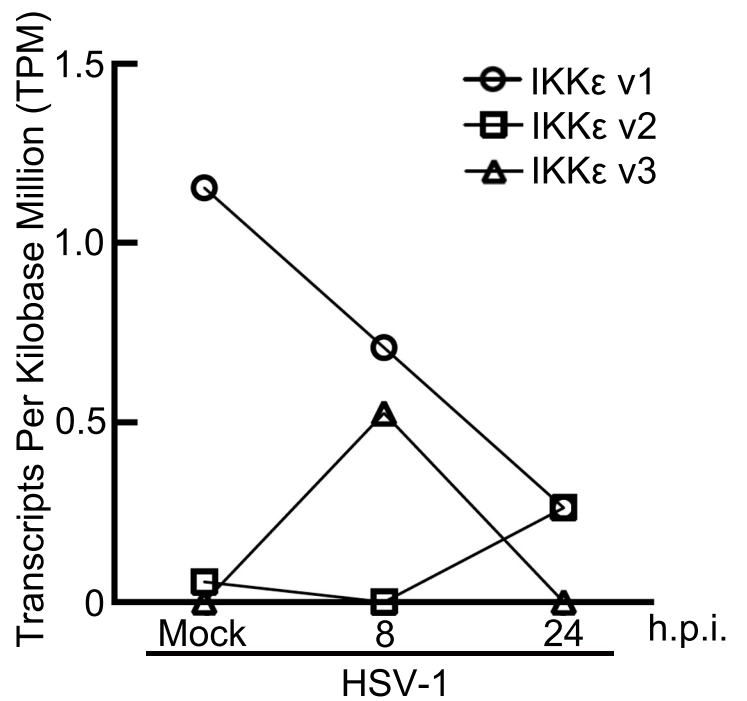
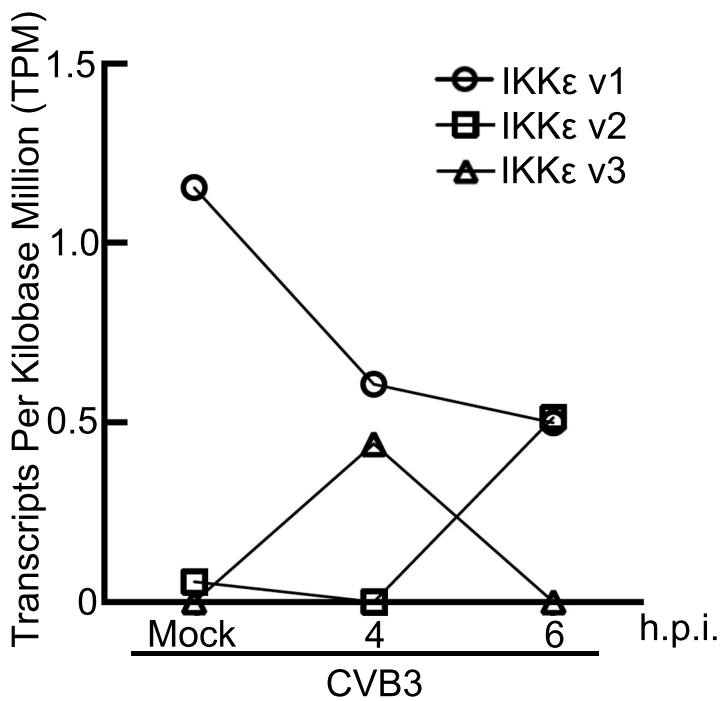
a



b



c



Supplementary Fig. 1: The pathway enriched by isoform switching genes in EV71 infection.

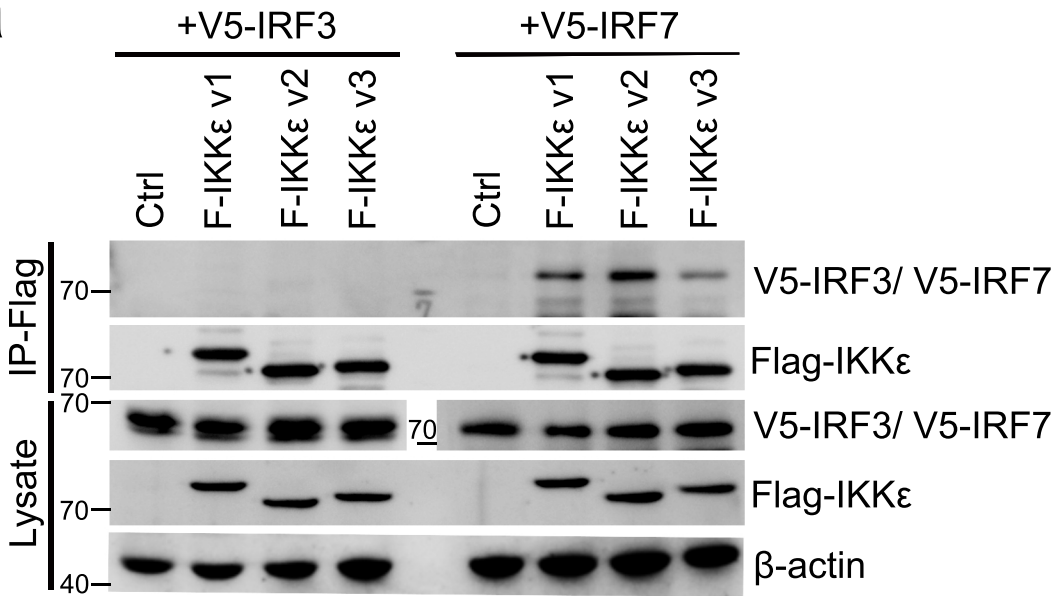
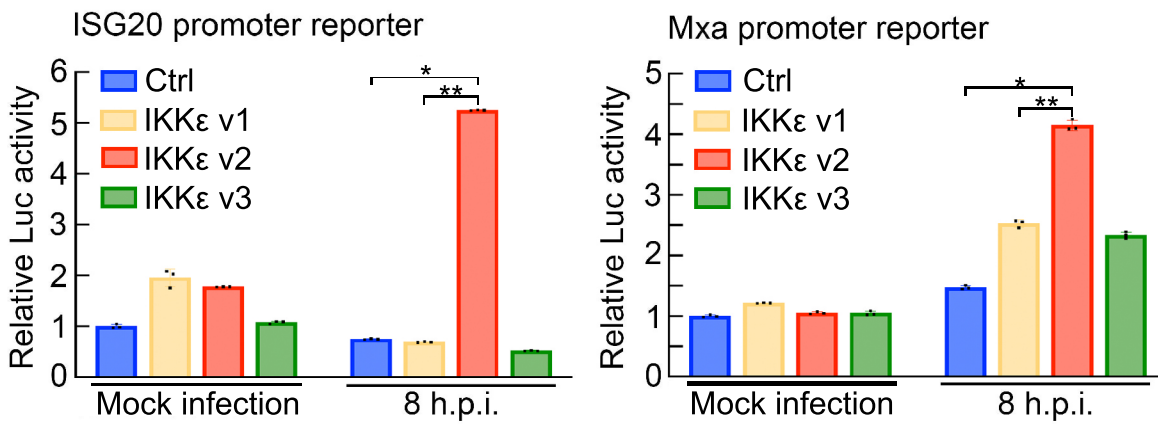
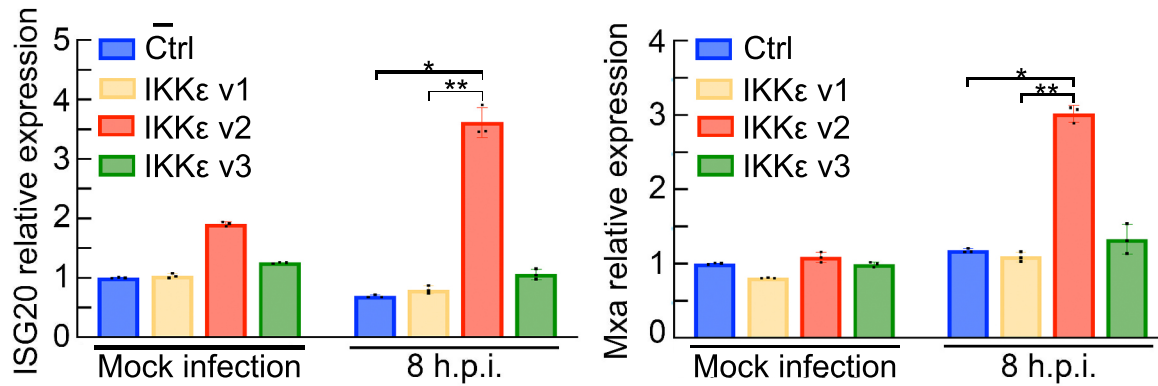
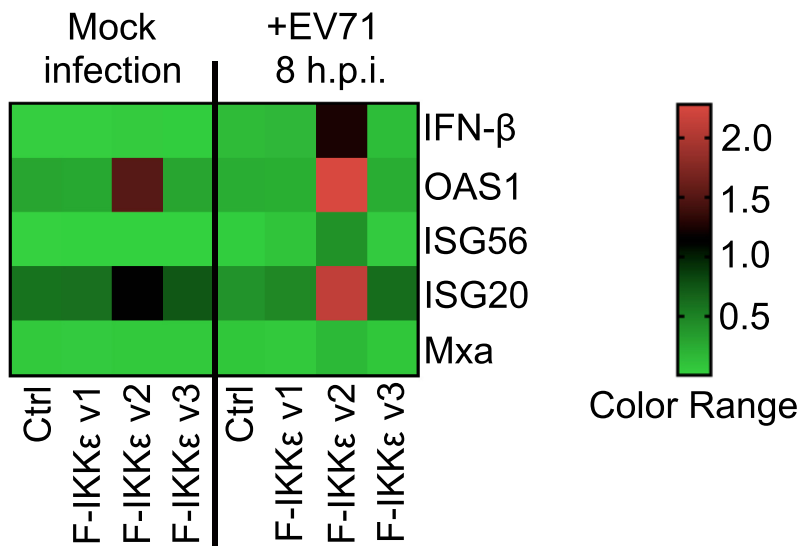
a. Toll-like receptor signaling pathway enriched in EV71 infection. The isoform switching genes identified in EV71 infection were adapted to DAVID Bioinformatics Resources analysis and the Toll-like receptor signaling pathway was ranked as top-one pathway with most significance (p value= 0.021). The illustrated pathway is a part of Toll-like receptor signaling pathway referenced from DAVID Bioinformatics Resources. The red circle and the gray circle represented the gene with and without isoform switching in EV71 infection, respectively.

b. Illustration of IKK ϵ isoforms. IKK ϵ gene possesses three isoforms: isoform 1 (IKK ϵ v1, NM_014002; NP_054721) contains full-length coding DNA sequence (CDS) while v2 (IKK ϵ v2, NM_001193322; NP_001180251) and v3 (IKK ϵ v3, NM_001193321; NP_001180250) truncates exon 20 and exon 3, respectively.

In protein level, IKK ϵ v2 is defected in coiled-coil domain functioned as protein-protein interaction and IKK ϵ v3 is trimmed in kinase domain contrast to IKK ϵ v1.

c. IKK ϵ isoform switching is measured by RNA-Seq in CVB3 and HSV-1 infected HeLa cells. IKK ϵ v2 was up-regulated while IKK ϵ v1 was down-

regulated in response to CVB3 (left panel) and HSV-1 (right panel) infection
determined by RNA-Seq.

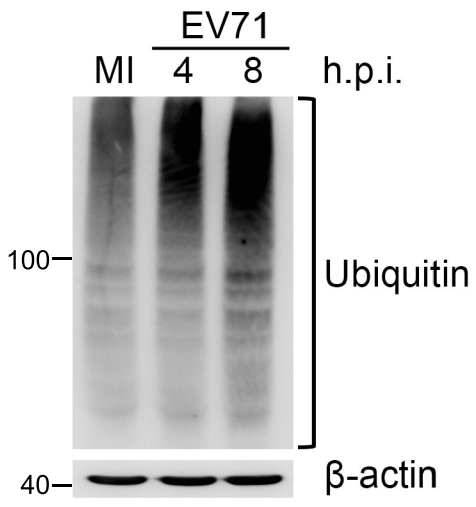
a**b****c****d**

Supplementary Fig. 2: IKK ϵ v2 transcriptionally induces ISGs**expressions in EV71 infection through IRF7.**

a. IKK ϵ preferentially binds to IRF7. Each Flag-IKK ϵ isoform was co-transfected with V5-IRF3 or V5-IRF7 in HEK293T cells. Flag-IKK ϵ was immunoprecipitated with anti-Flag beads. V5-IRF3 (left panel) and V5-IRF7 (right panel) were detected by anti-V5 antibody.

b, c. IKK ϵ v2 up-regulates promoter activities and expressions of MxA and ISG20. Each IKK ϵ isoform was ectopically expressed in RD cells accompanied with the indicated ISG promoter followed by EV71 infection. The promoter activities (b) and ISGs expressions (c) were measured at indicated h.p.i.. The data are normalized with vector control (Ctrl) in mock infection group. * and ** represent p value <0.05 as compared with Ctrl group and IKK ϵ v1 group, respectively. All data presented are mean \pm SD (n=3).

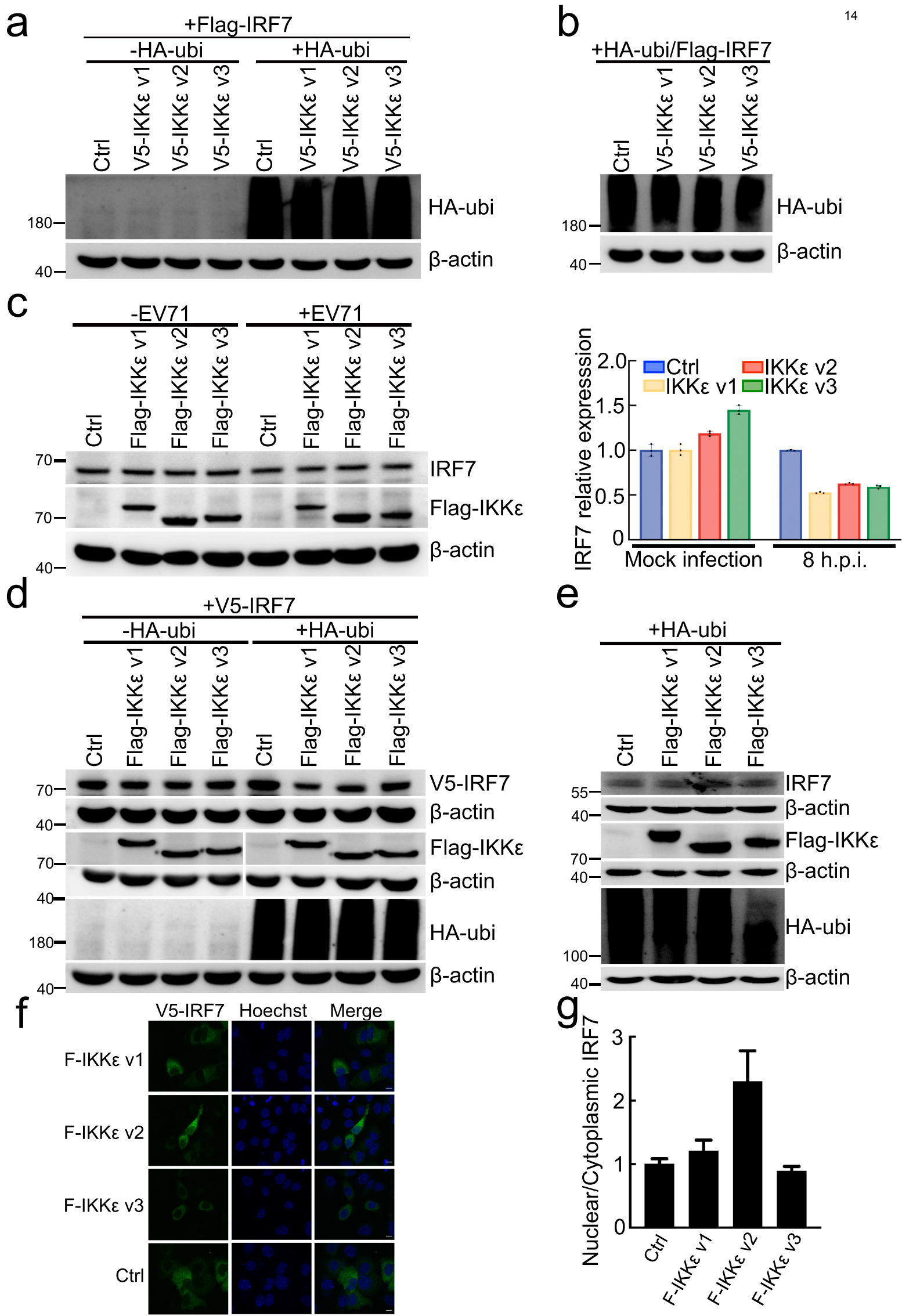
d. The heat map of ISGs expressions in ectopically IKK ϵ isoform-expressed RD cells, referred to Fig. 2b, 2d, S2a and S2b. The color scale corresponds to the mean for each gene.



Supplementary Fig. 3: EV71 infection promotes ubiquitination.

RD cells were infected with EV71 at 10 m.o.i. and harvested at indicated h.p.i..

The total lysates were adapted to immunoblot analysis with anti-ubiquitin antibody. β -actin was served as an internal control. MI: mock infection



Supplementary Fig. 4: IKK ϵ v2 increases IRF7 phosphorylation and IRF7 translocation in the presence of ubiquitin.

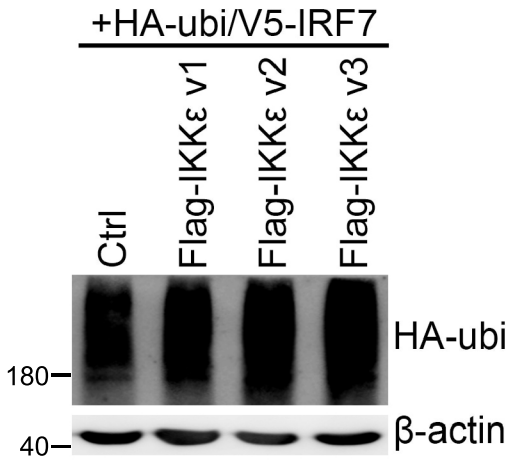
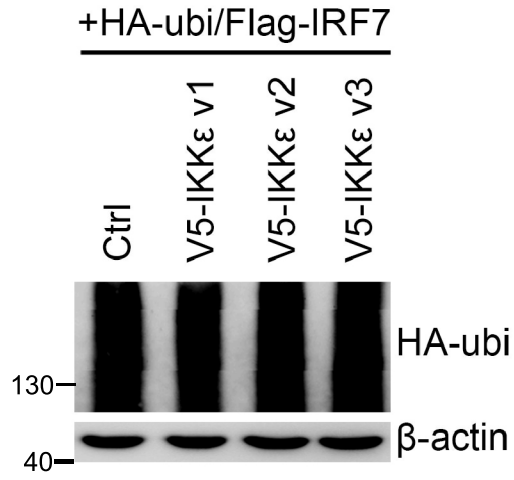
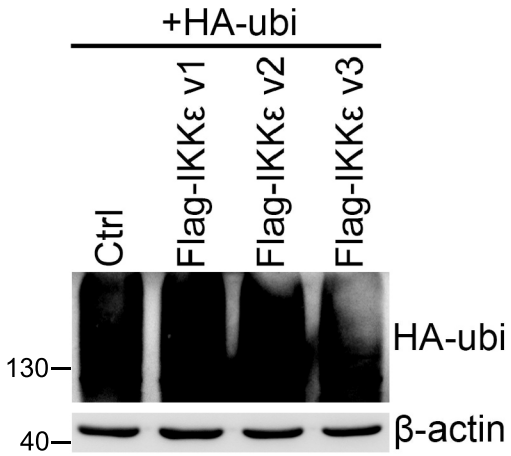
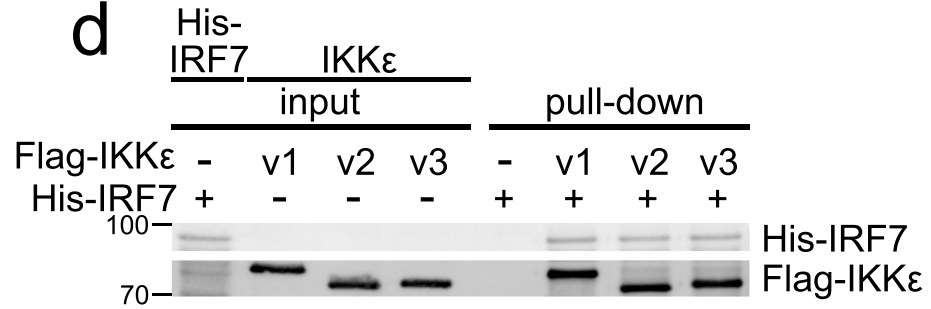
a, b. The expression levels of HA-ubiquitin referred to Fig. 3a, 3b. HEK293 or HeLa cells were transfected with indicated plasmids and total lysates were loaded to perform immunoblot with indicated antibodies. β -actin served as an internal control.

c. IRF7 mRNA and protein were not induced by IKK ϵ v2. RD cells were ectopically expressed Flag- IKK ϵ followed by EV71 infection. Total lysates were loaded to perform immunoblot with indicated antibodies. β -actin served as an internal control (left panel). RNA expression of IRF7 was measured by quantitative real-time PCR and normalized with Ctrl in mock infection group (right panel). All data presented are mean \pm SD (n=3).

d, e. The expression levels of HA-ubiquitin referred to Fig. 3d, 3e. HeLa cells were transfected with indicated plasmids and total lysates were loaded to perform immunoblot with indicated antibodies. β -actin served as an internal control.

f. IKK ϵ v2 facilitates IRF7 translocation by immunofluorescence analysis. The immunofluorescence images of cells treated with the same condition used in Figure 3d. Scale bar: 10um

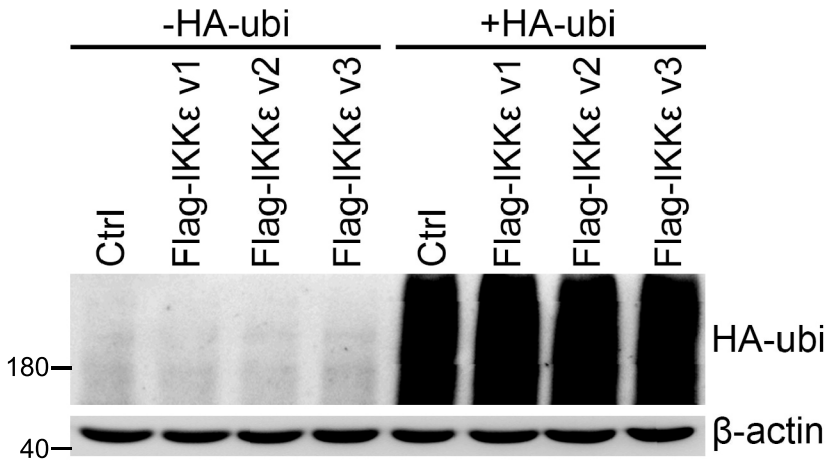
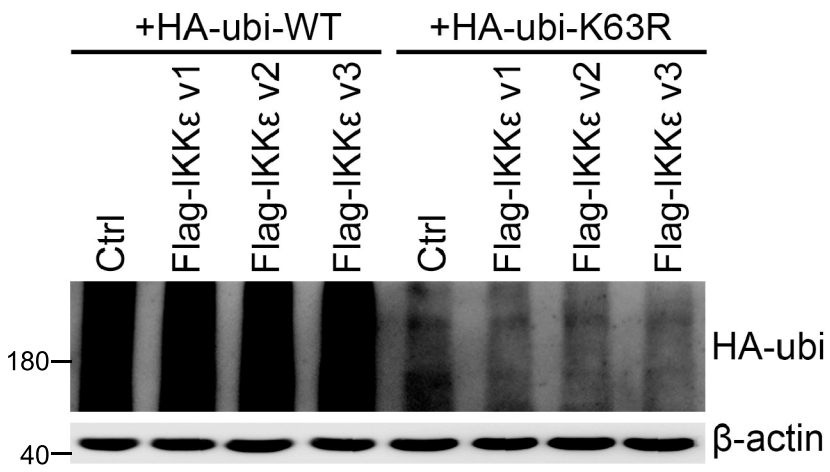
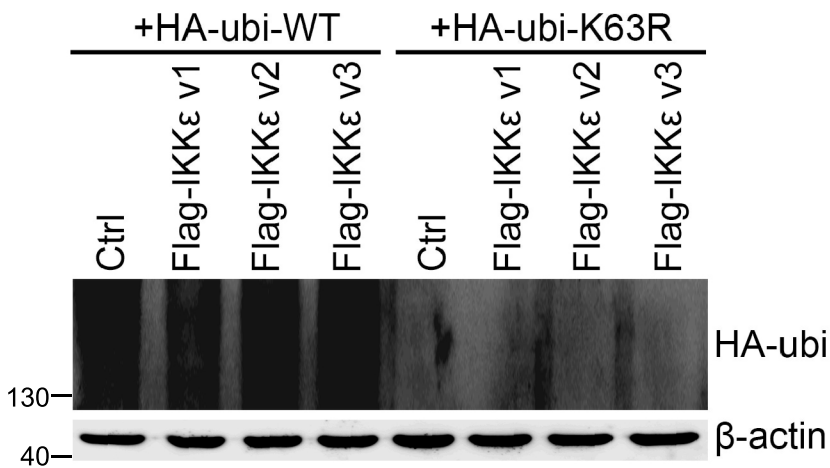
g. The ratio of nuclear IRF7 to cytoplasmic IRF7 in Supplementary Figure 3f. The ratios of F- IKK ϵ isoforms were normalized to that of Ctrl.

a**b****c****d**

Supplementary Fig. 5: The interaction of IKK isoforms and IRF7.

a-c. The expression levels of HA-ubiquitin, referred to Fig. 4. HEK293 cells were transfected with indicated expressing plasmids and total lysates were loaded to perform immunoblot with anti-HA antibody. β -actin served as an internal control.

d. The direct interaction of IRF7 and IKK ϵ isoforms. Each Flag-IKK ϵ isoform was purified by anti-Flag beads from Flag-IKK ϵ -expressed HEK293T cells and incubated with 1 μ g of His-IRF7, which was purified from *E. coli*. for 2 hours at 4°C. After washing for three times, the bound His-IRF7 was analyzed by Western blotting with anti-His antibody.

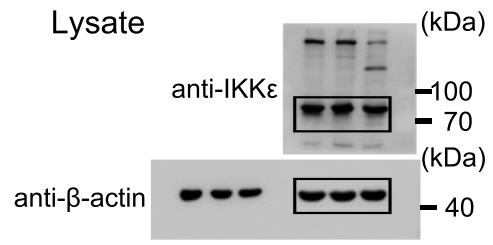
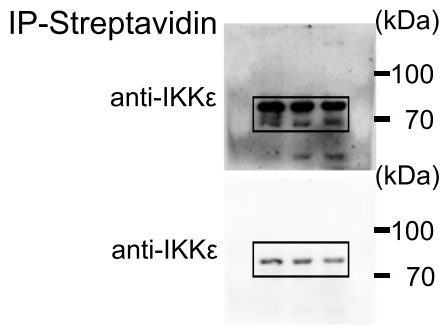
a**b****c**

Supplementary Fig. 6: The expression levels of HA-ubiquitin, referred to Figure 5.

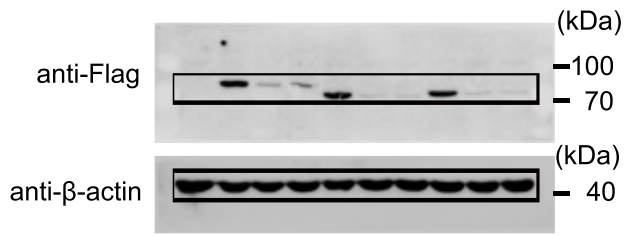
a-c. HEK293 cells were transfected with indicated expressing plasmids and total lysates were loaded to perform immunoblot with anti-HA antibody. β -actin served as an internal control.

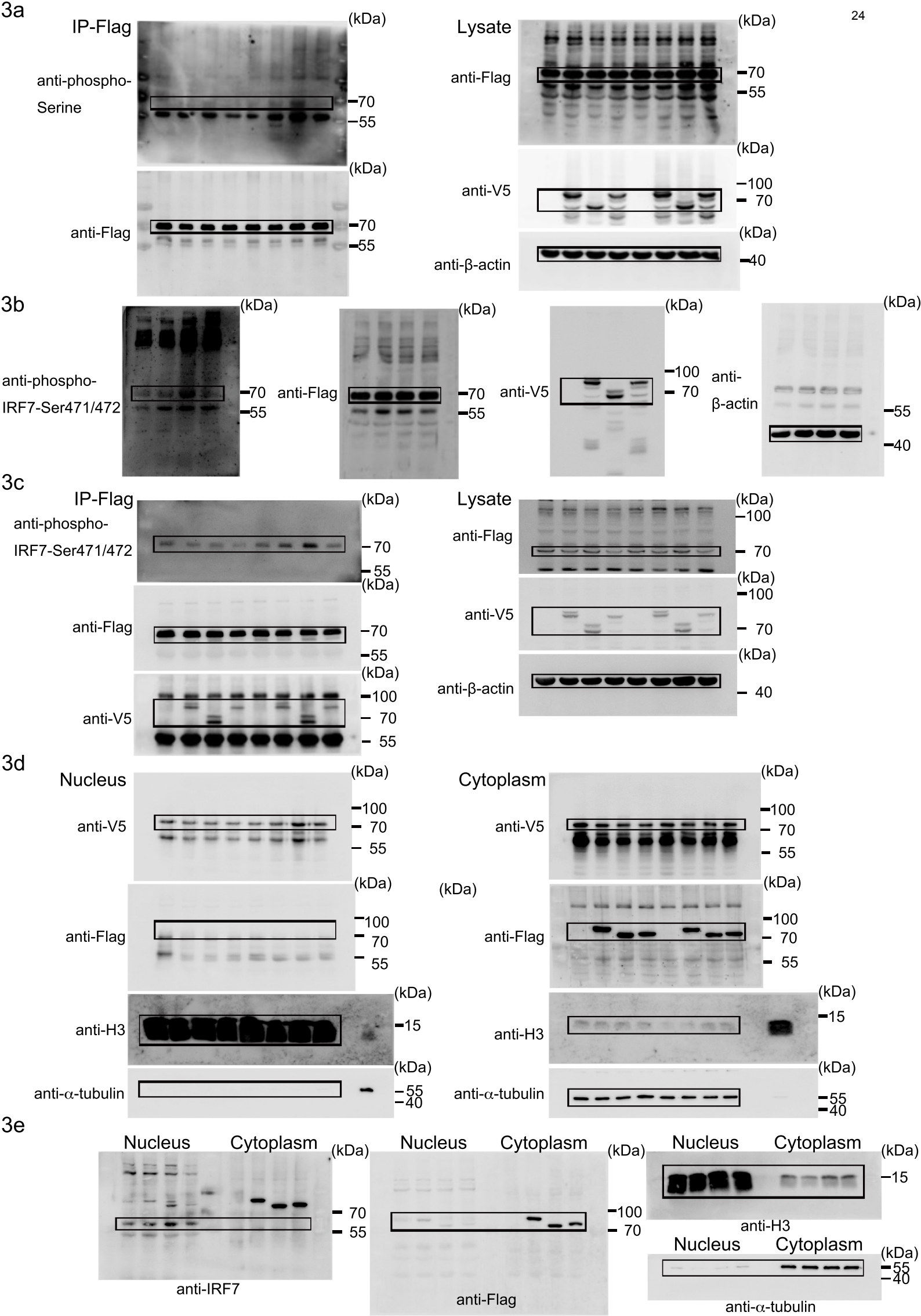
Supplementary Figure 7-11: Full-scan images of representative Western blots, referred to Figure 1-5

1c

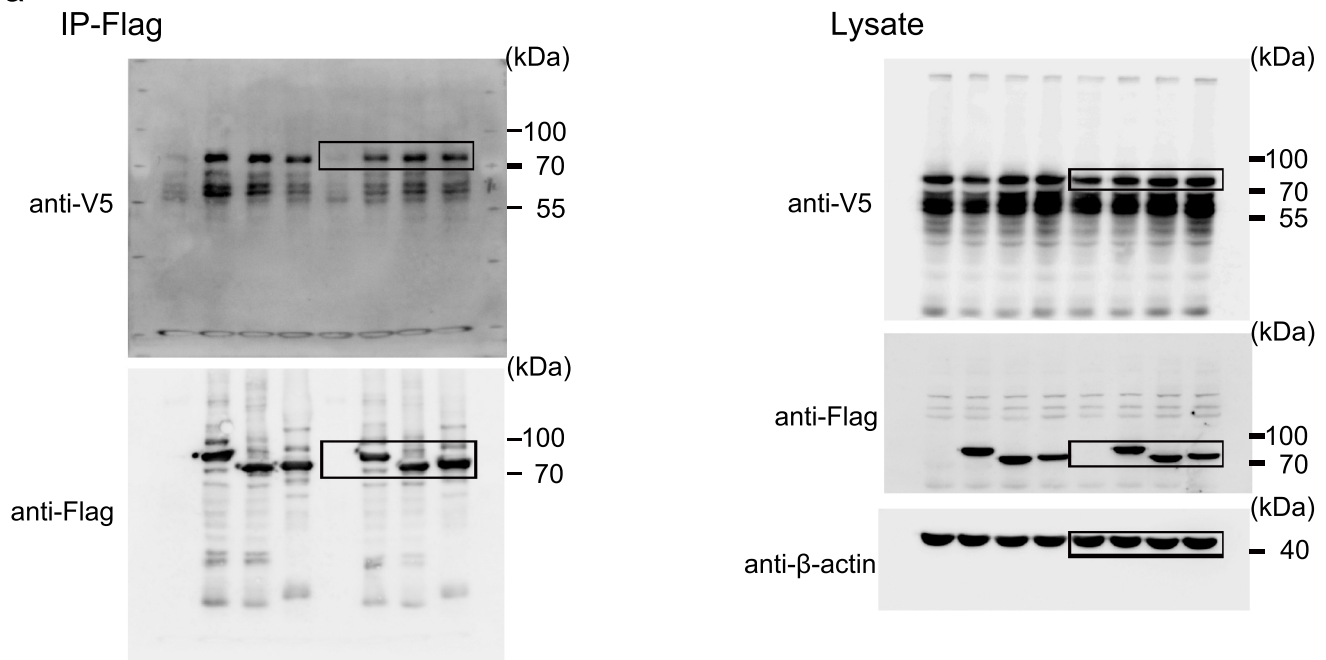


2f

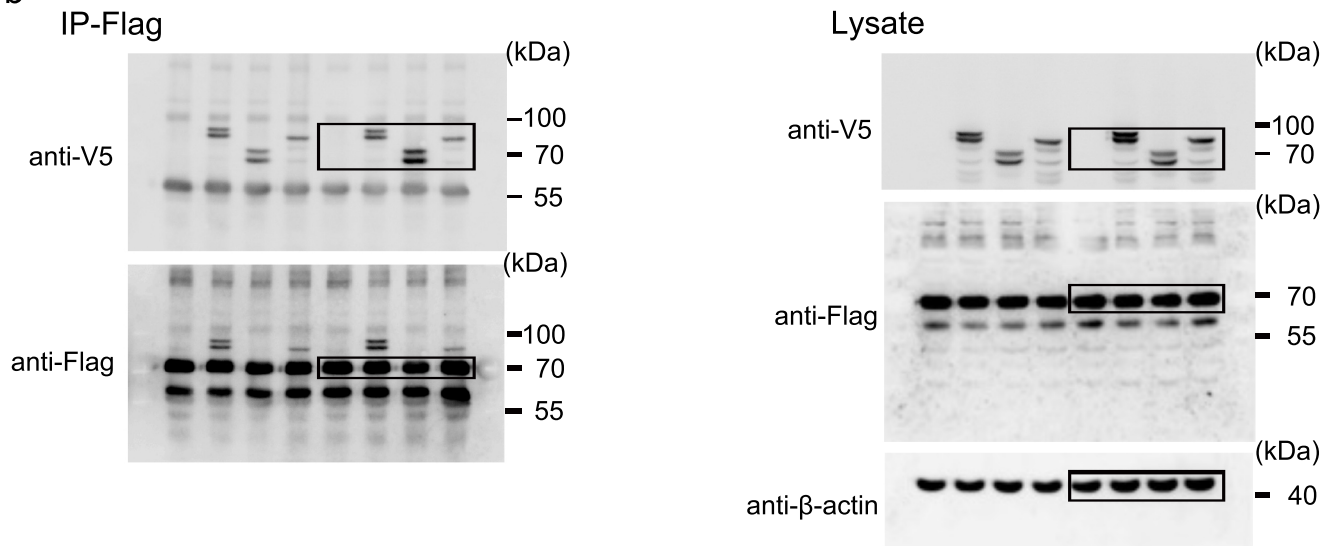




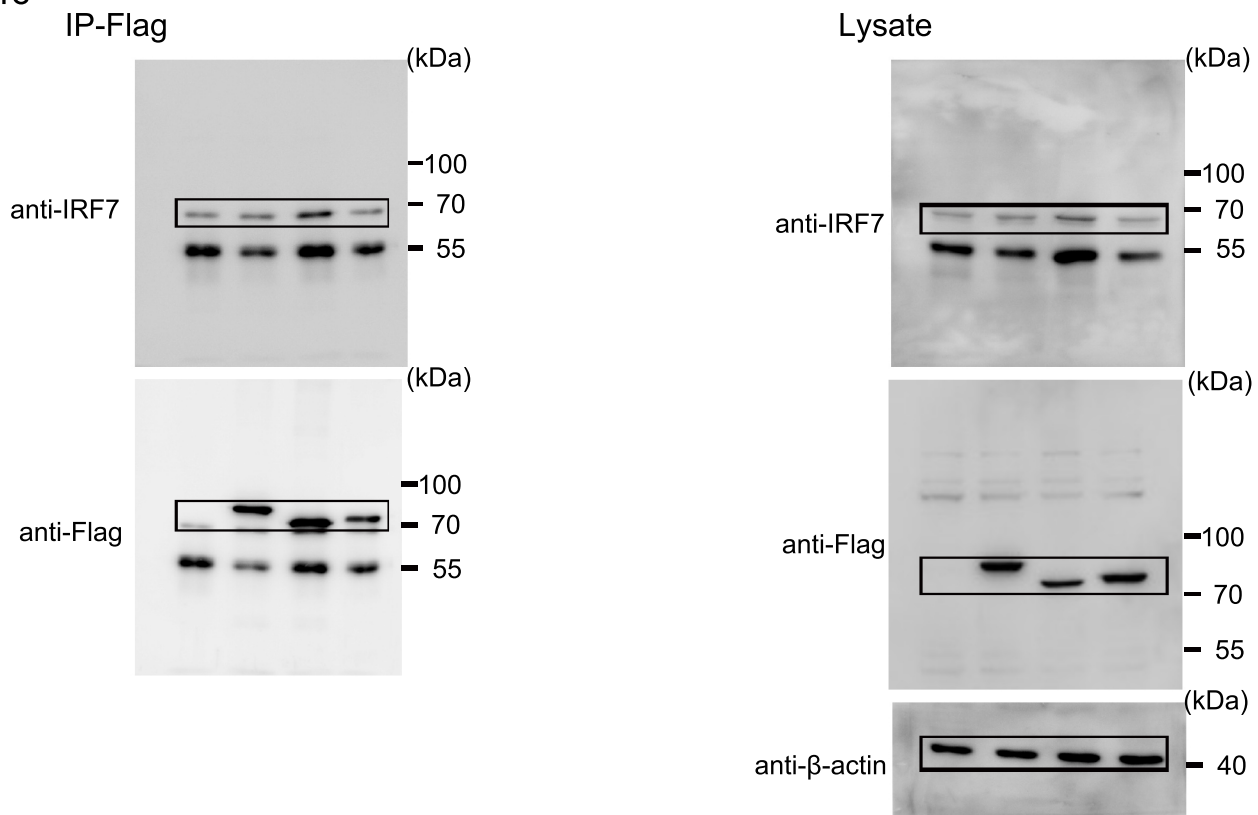
4a



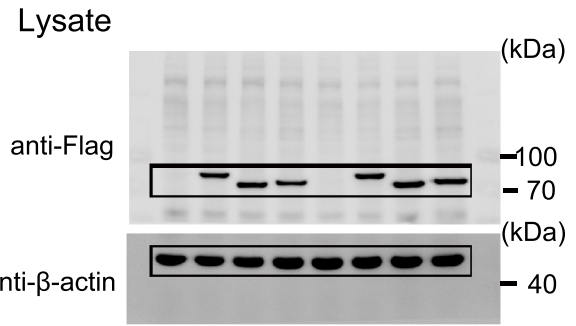
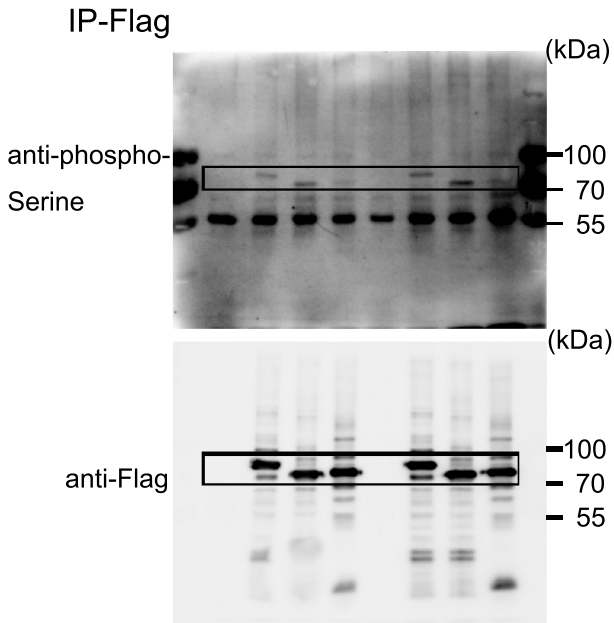
4b



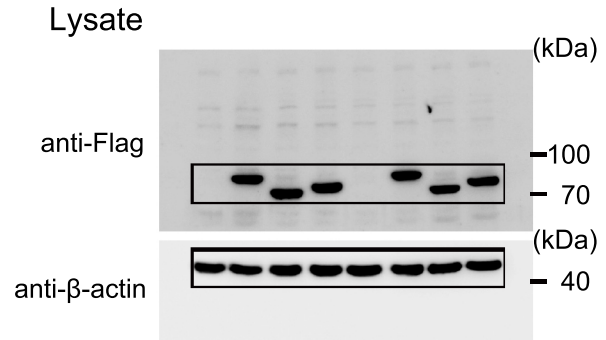
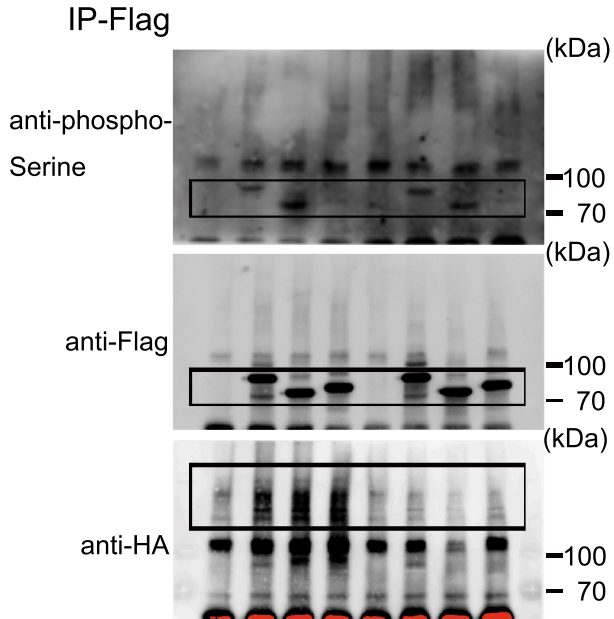
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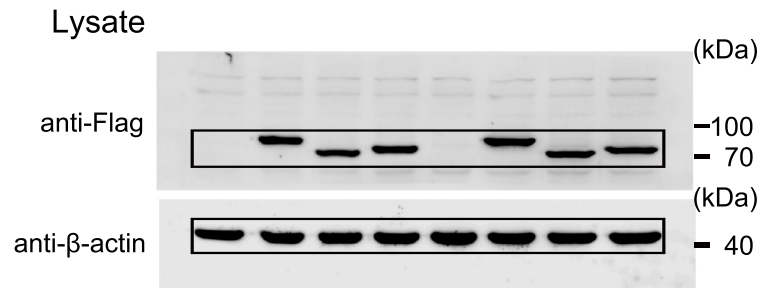
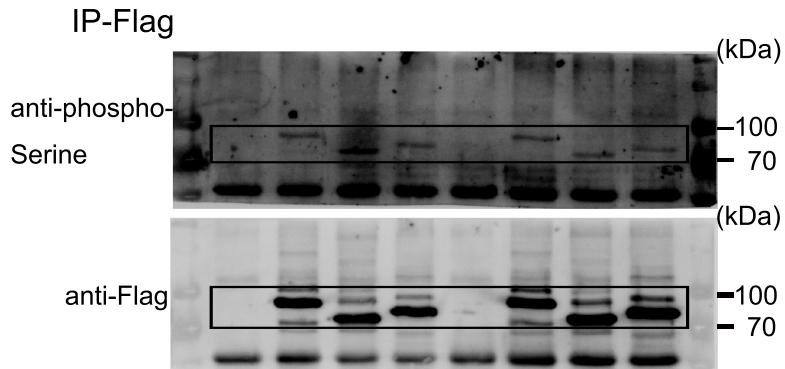
5a



5b

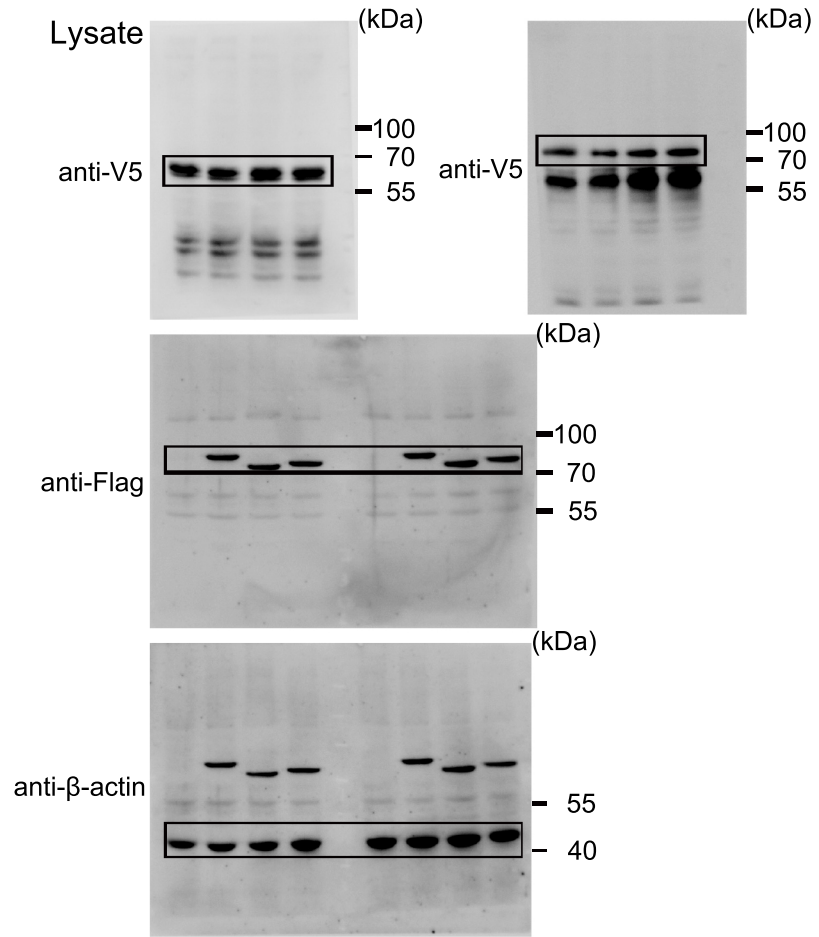
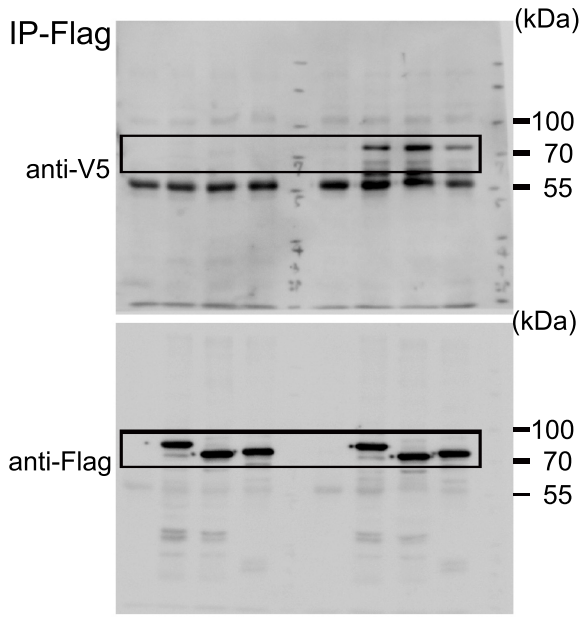


5c

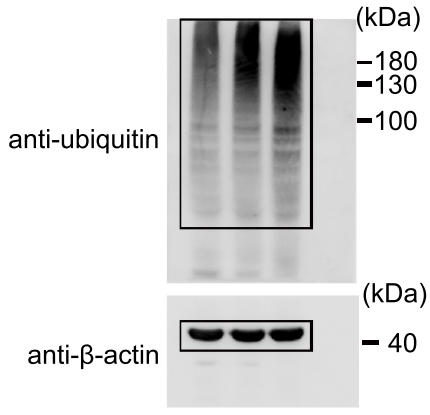


Supplementary Figure 12-16: Full-scan images of representative Western blots, referred to Supplementary Figure 2-6

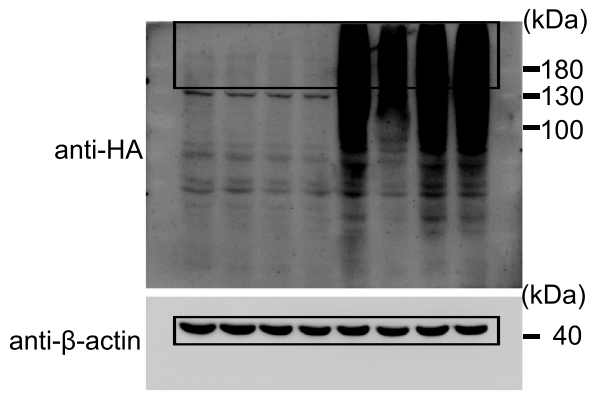
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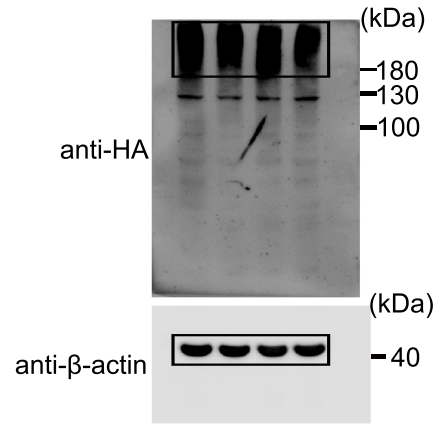
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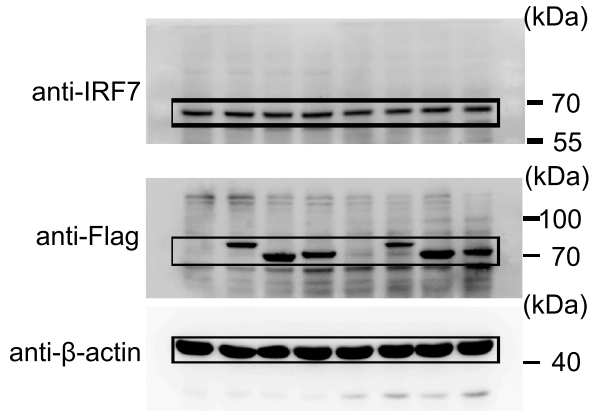
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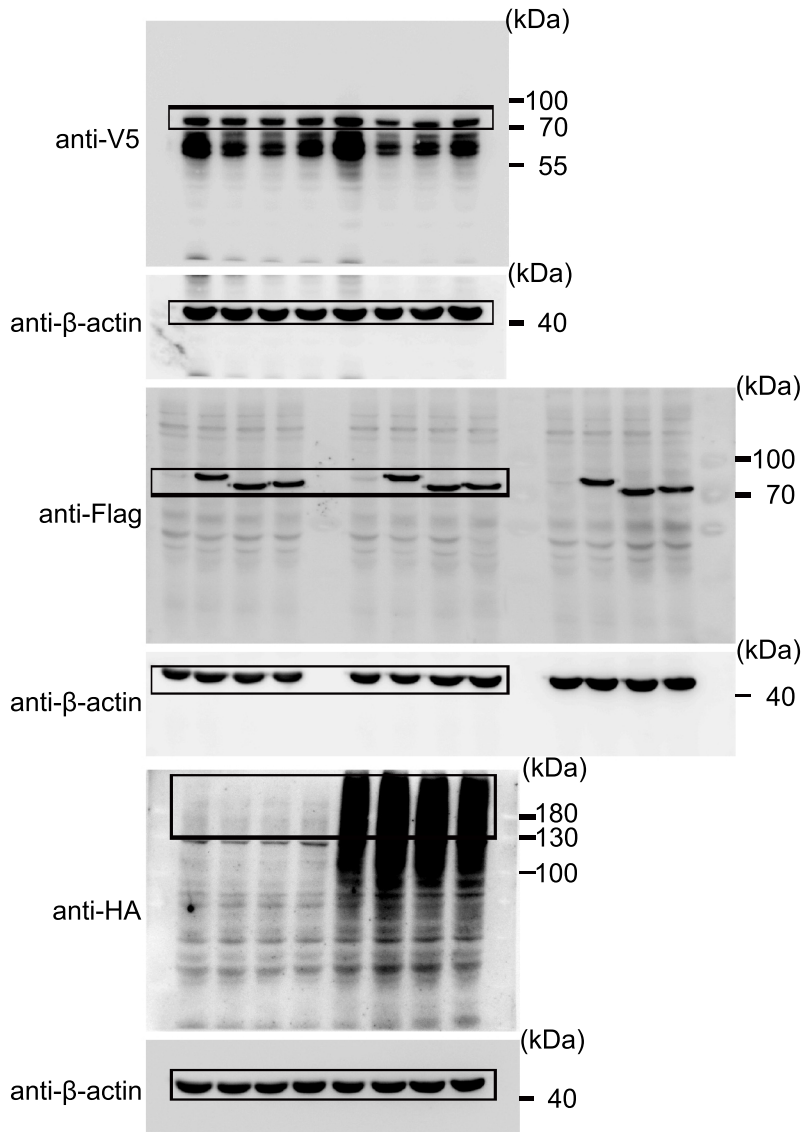
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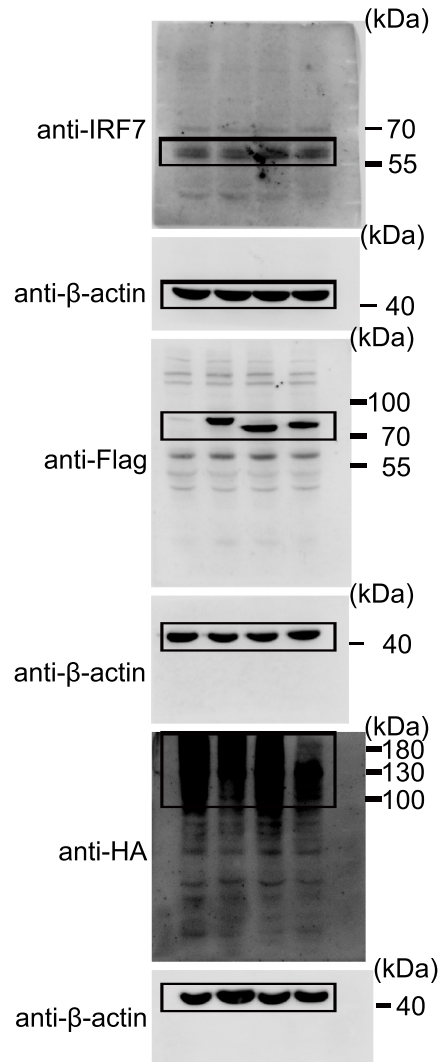
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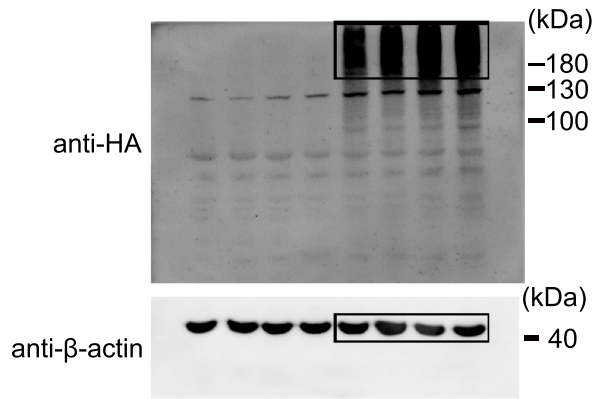
S4d



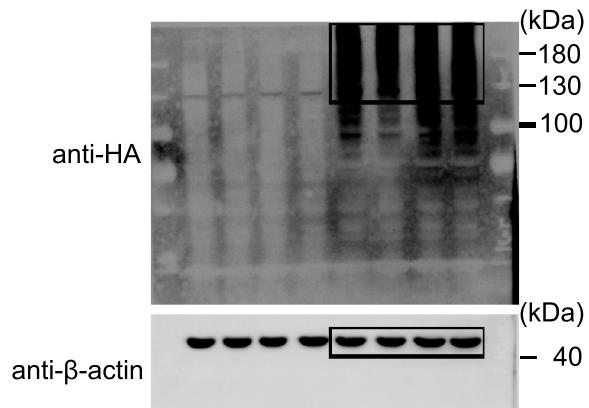
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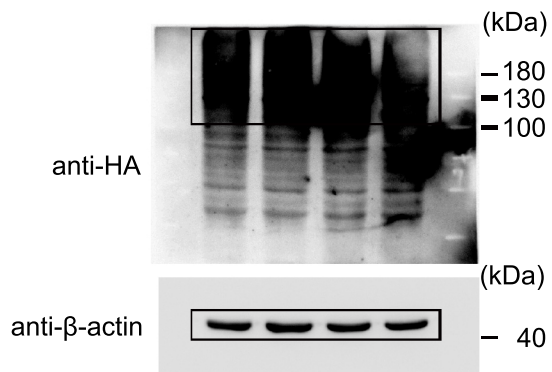
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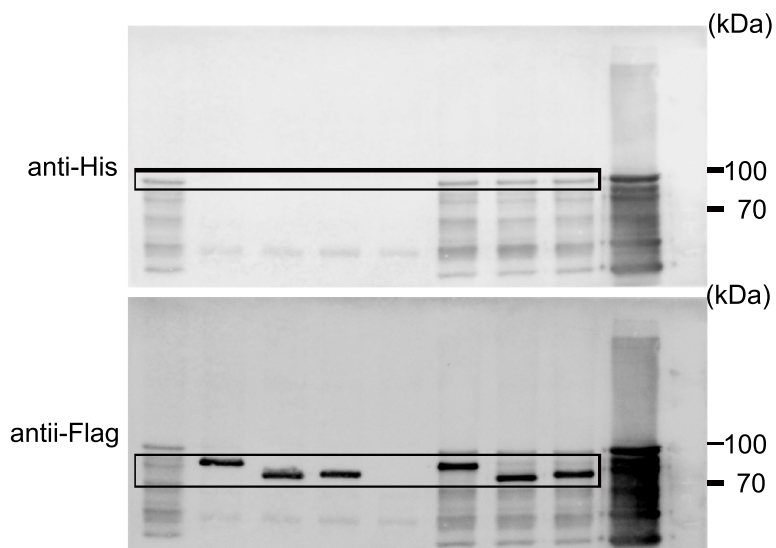
S5b



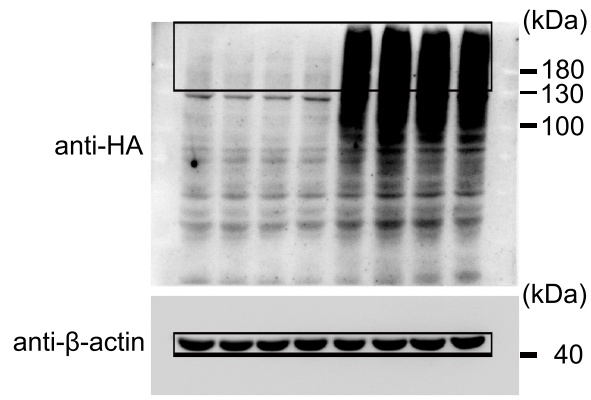
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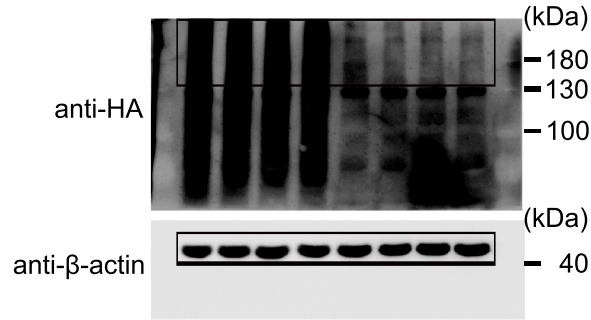
S5d



S6a



S6b



S6c

