

## **SUPPLEMENTARY MATERIAL**

### **NF- $\kappa$ B promotes leaky expression of adenovirus genes in a replication-incompetent adenovirus vector**

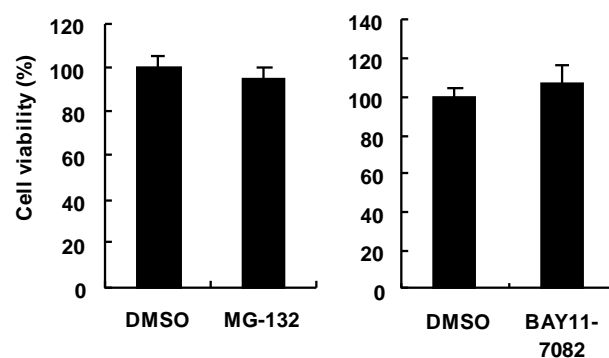
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#### **Supplementary Materials and Methods**

##### ***Cell viability assay***

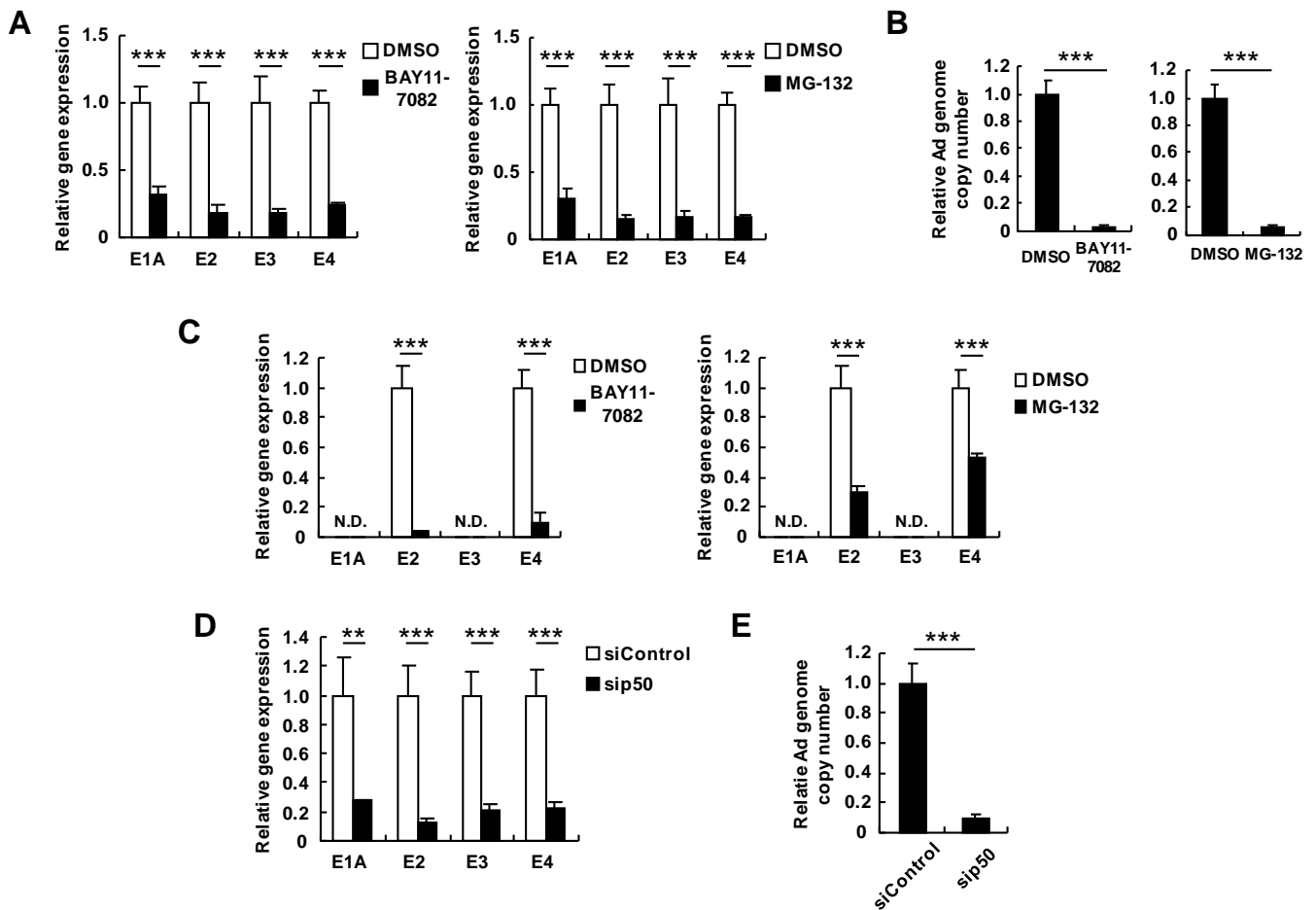
Cells were treated with BAY11-7082 and MG-132. After 24-h incubation, cell viabilities were determined by staining with AlamarBlue (Life Technologies, Carlsbad, CA) according to the manufacturer's instructions. Cell viabilities of DMSO-treated cells were normalized to 100%.

## Supplementary Figures



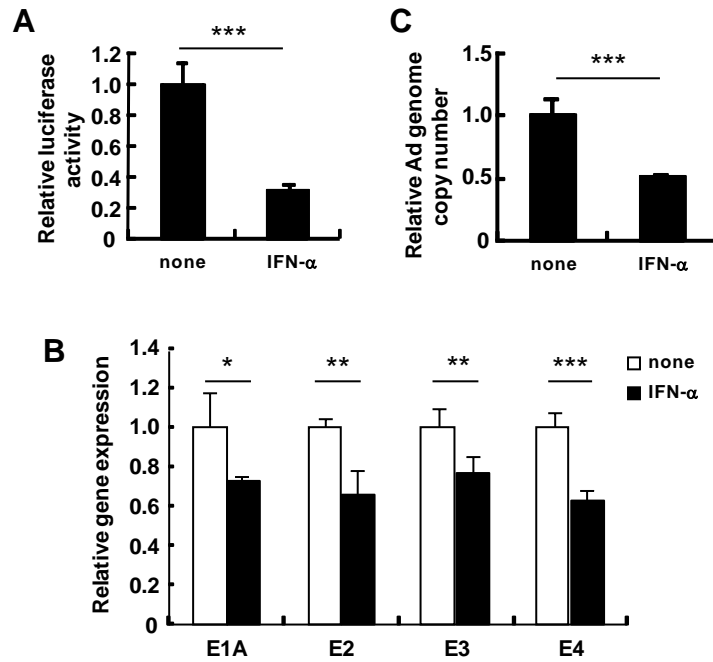
**Figure S1 Cell viability following treatment of MG-132 and BAY11-7082.**

HeLa cells were treated with MG-132 and BAY11-7082 at 2.5 mM and 10 mM, respectively. After 24-h incubation, cell viability was determined by AlamarBlue assay. Cell viabilities of DMSO-treated cells were normalized to 100%. These data are expressed as the means  $\pm$  S.D. (n=4).



**Figure S2 Suppression of Ad early gene expression and Ad replication by inhibition of NF- $\kappa$ B.**

(A-C) H1299 cells were pre-treated with BAY11-7082 and MG-132 at 10 mM and 2.5 mM, respectively, for 1 h, followed by infection with WT-Ad or Adv-CMVLuc at 100 VP/cell. After 12-h incubation, the E1A, E2, E3, and E4 mRNA levels in the cells were determined by quantitative RT-PCR (A, C). After 24-h incubation, Ad genome copy numbers in the cells were determined by real-time PCR (B). (D, E) H1299 cells were transfected with sip50 at 50 nM, followed by infection with WT-Ad at 100 VP/cell. After 12-h incubation, Ad gene mRNA levels in the cells were similarly determined (D). After 24-h incubation, Ad genome copy numbers in the cells were similarly determined (E). These data are expressed as the means  $\pm$  S.D. (n=3-4). \*\*p<0.01, \*\*\*p<0.001.



**Figure S3 Suppression of Ad early gene expression and replication by IFN- $\alpha$  stimulation.**

(A) HeLa cells were transfected with pNF- $\kappa$ B-Luc, followed by treatment with IFN- $\alpha$  at 1000 U/ml. After 24-h incubation, luciferase activity was determined. The data show FLuc activity normalized by RLuc activity. (B) HeLa cells were pre-treated with IFN- $\alpha$  at 1000 U/ml for 5 h, followed by infection with WT-Ad at 100 VP/cell. After 12-h incubation, the E1A, E2, E3, and E4 mRNA levels in the cells were determined by quantitative RT-PCR. (C) HeLa cells were pre-treated with IFN- $\alpha$  at 1000 U/ml for 5 h, followed by infection with WT-Ad at 100 VP/cell. After 24-h incubation, Ad genome copy numbers in the cells were determined by quantitative PCR. These data are expressed as the means  $\pm$  S.D. (n=3-4). \*p<0.05, \*\*p<0.01, \*\*\*p<0.001.

**Table S1****The oligonucleotides and primers used in this study**

No	name	sequence
1	E1Ap-F	atactcgagcatcatcaataatactattttggattga
2	E1Ap-R	tataagcttgctggagcggctcggag
3	E1Bp-F	atactcgagggtctagagaatgcaatagtag
4	E1Bp-R	ataaagcttaaccaagattgcccacgg
5	E2p-F	atactcgagtaggattgcctgacgaggcg
6	E2p-R	ataaagcttactgcgcgctgactcttaagg
7	E3p-F	atactcgaggcagctgcctgtatcacaaa
8	E3p-R	ataaagcttagctgaatactcgcctct
9	E4p-F	gcgaagcttcagtcagcctaccagtaaaaaag
10	E4p-R	gcgctcgagcatcatcaataatactattttgg
11	E2-del2-F	atactcgaggctgtaactccacatgtag
12	E2-del2-R	atggtggctttaccaacag
13	E2-del3-S	tcgagctggagatgacgtagtttcgcgctaaattgagaaagggcgcgaaactagtc
14	E2-del3-AS	ttaagactagtttcgcgccctttctcaaatttaagcgcgaaaactacgcatctccagc
15	E2-del2.1-F	gtaactccacatgtagggcgtcaattgctcataatggcgctg
16	E2-del2.1-R	cagcgccattatgagcaattgacgcctacatgtggagtac
17	DNikBa-F	tgtctagacagccatgtttcagccagc
18	DNikBa-R	aagcggcgcgtataatgtcagacgctggcc
19	E1A-F	tccggtccttctaacacacctc
20	E1A-R	acggcaactggttaatggg
21	E2-F	cactacggtgcgagtgcaa
22	E2-R	ggtagctgcctcccaaaaag
23	E3-F	aacacctggtccactgtcgc
24	E3-R	agctcggagaggttctctcgtag
25	E4-F	gggatcgtctacctctttga
26	E4-R	gggcagcagcggatgat
27	pIX-F	gcccgcgggatttg
28	pIX-R	cgggaagctgcactgctt
29	p50-F	aacagagaggatttcgtttccg
30	p50-R	ttgacctgagggttaagacttct
31	hGAPDH-F	ggtggtctcctctgactcaaca
32	hGAPDH-R	gtggtcgttgagggaatg

33	mGAPDH-F	caatgtgtccgtcgtggatct
34	mGAPDH-R	gtcctcagtgtagcccaagatg
35	ChIP E2/E3-F	agcgcgaaaactacgcatc
36	ChIP E2/E3-R	tcccattgtggctggtaac