

Supplemental Data

Figure S1. Related to Figure 1: Silencing USP7 Decreases NEK2-induced

Bortezomib Resistance

OCI-MY5 cells transduced with EV, NEK2-OE or NEK2-OE + USP7-shRNA were treated with DOX to suppress USP7 expression in NEK2-OE cells. After 72 hours, cells were treated with bortezomib (5nM) for further 24 hours and cell viability was measured using trypan blue stain.

Figure S2. Related to Figure 2: USP7 Deubiquitinizes NEK2 Protein

(A) H1299 cells were transfected with USP7-shRNA, after 72-hour induction with DOX, followed by protein extraction. Western blot were performed using anti-USP7, NEK2 and β -actin antibodies. (B) ARP1 cells were treated with P5091 overnight at 16 μ M. Cells were lysed, and protein extract was analyzed by western blot using USP7, USP47, NEK2, and GAPDH antibodies. (C) ARP1 cells were treated with P5091 overnight at 16 μ M. Cells were lysed, total RNA was extracted and analyzed for NEK2 mRNA expression by qRT-PCR (NS: No Significance). (D & E) OCI-MY5 (D) and H1299 (E) cells were treated with P5091 at different doses or a combination with the proteasome inhibitor MG132 for 5 hours. NEK2 protein levels were analyzed by western blot using NEK2 and β -actin antibodies. (F) Input controls for immunoprecipitation of Figure 2H. H1299 cells were co-transfected with HA-UB and NEK2-OE vectors with or without a USP7-OE plasmids. Cells were lysed in the presence of NEM and protein extracts were analyzed by western blot using NEK2, USP7, and β -actin antibodies.

Figure S3. Related to Figure 3: Overexpression of NEK2 increases phosphorylated p-65 in myeloma cell lines. ARP1 and OCI-MY5 myeloma cells overexpressing NEK2 were lysed. NEK2 and p65-S536 phosphorylation levels were analyzed between NEK2-OE and EV cells by western blot.

Figure S4. Related to Figure 4: NEK2 Activates NF- κ B signaling via AKT

(A) H1299 cells transfected with EV or NEK2-OE were lysed and analyzed by western blot using NEK2, P-I κ B α , and GAPDH antibodies. (B) H1299 cells transfected with EV or NEK2-OE were treated with vehicle or MK-2206 2HCl for 30 minutes. Protein was extracted and analyzed by western blot using P-p65-S536, and β -actin antibodies. (C) H1299 cells transfected with NEK2-shRNA or Scramble were induced with DOX for 48 hours and then treated with tautomycin for another 24 hours. Cells were lysed and analyzed by western blot using NEK2, P-p65-S536, P-PP1 α , and P-AKT-S473 antibodies.

Figure S5. Related to Figure 7: INH1 and P5091 Overcome NEK2-induced Bortezomib Resistance

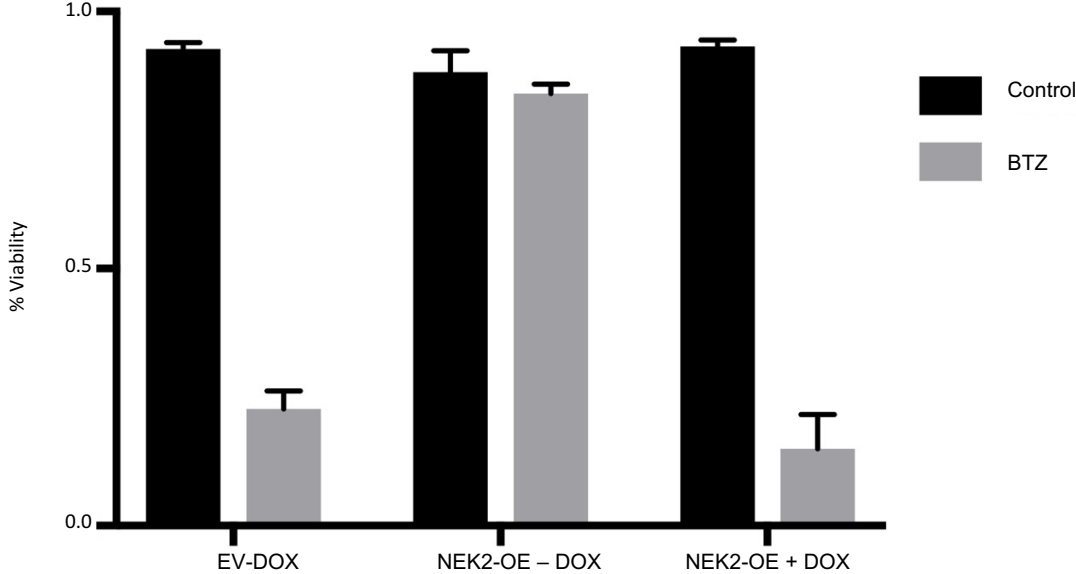
Mouse weight was calculated at the starting and ending of the injection of NEK2-OE ARP1 cells expressing luciferase into NOD-Rag1^{null} mice. After one-week injection of NEK2-OE cells, mice were treated with (1) vehicle, (2) BTZ (3mg/kg, IP, 2 times/week), (3) INH1 (100mg/kg, IP, 3 times/week), (4) P5091 (10mg/kg, IV, 2 times/week), (5) P5091 + BTZ, and (6) INH1 + BTZ Mice for 28 days.

Table S1. Related to Figure 1: Protein List Identified by TAP-MS Bound to NEK2.

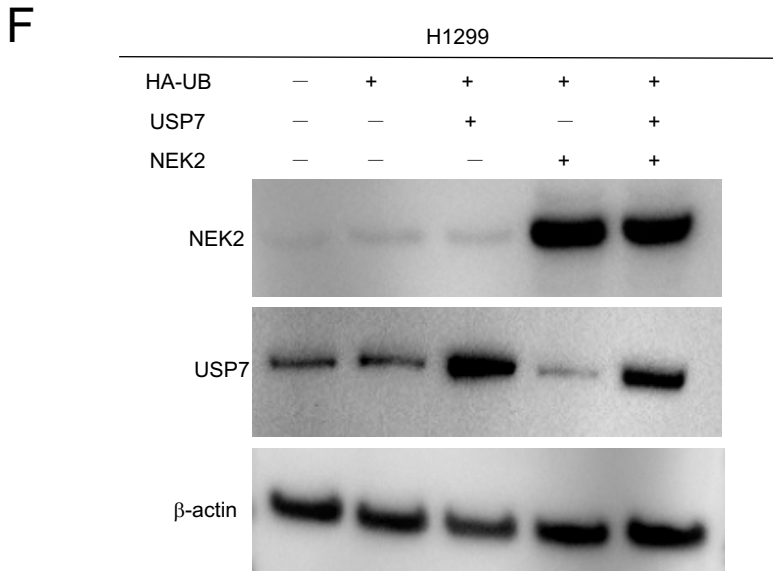
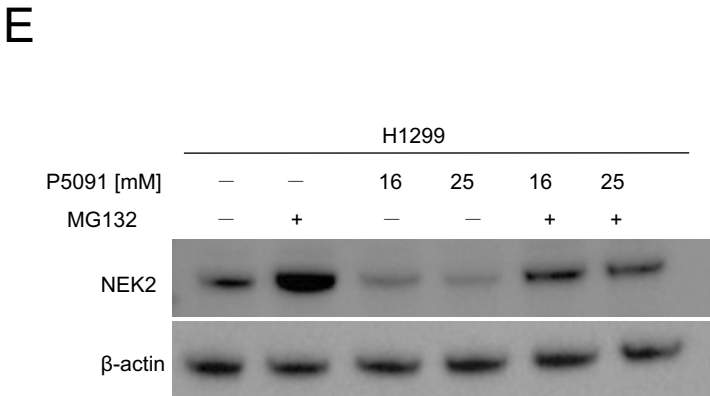
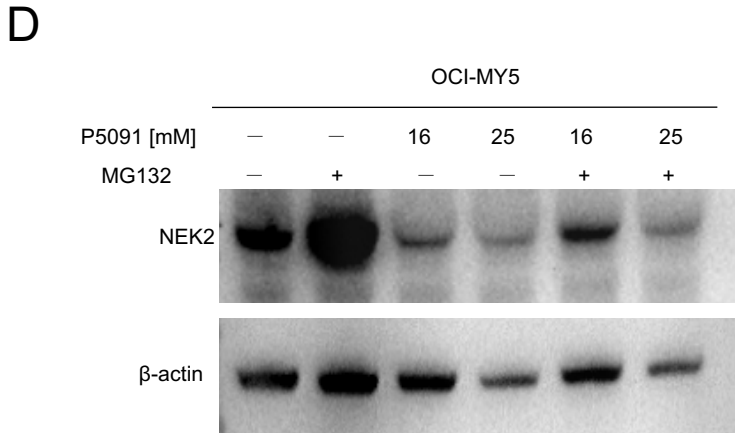
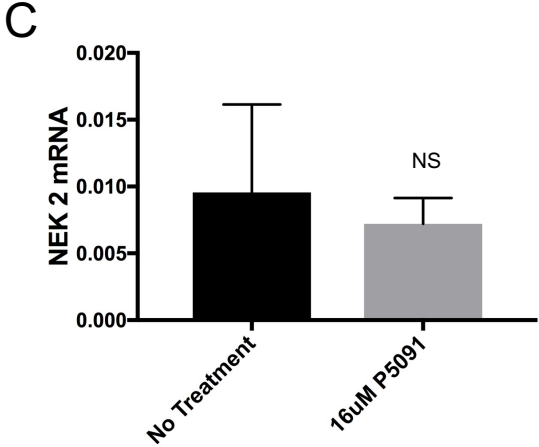
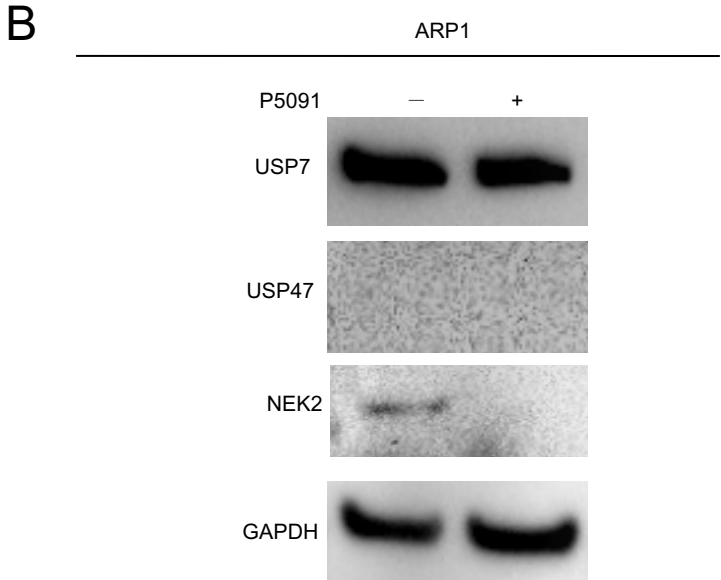
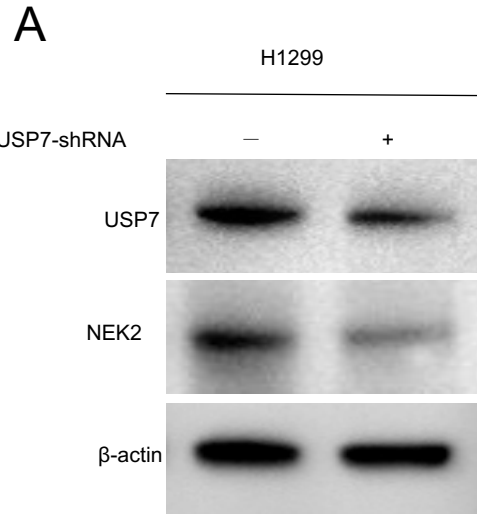
Tables S2. Related to Figure 5: p65 Binding Sites to Direct Target Genes

Regulated by NEK2 Identified by ENCODE at UCSC.

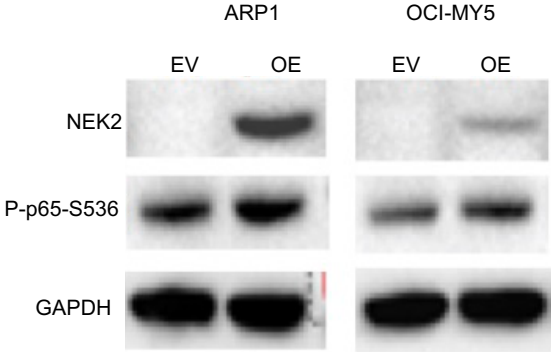
Supplemental Figure 1



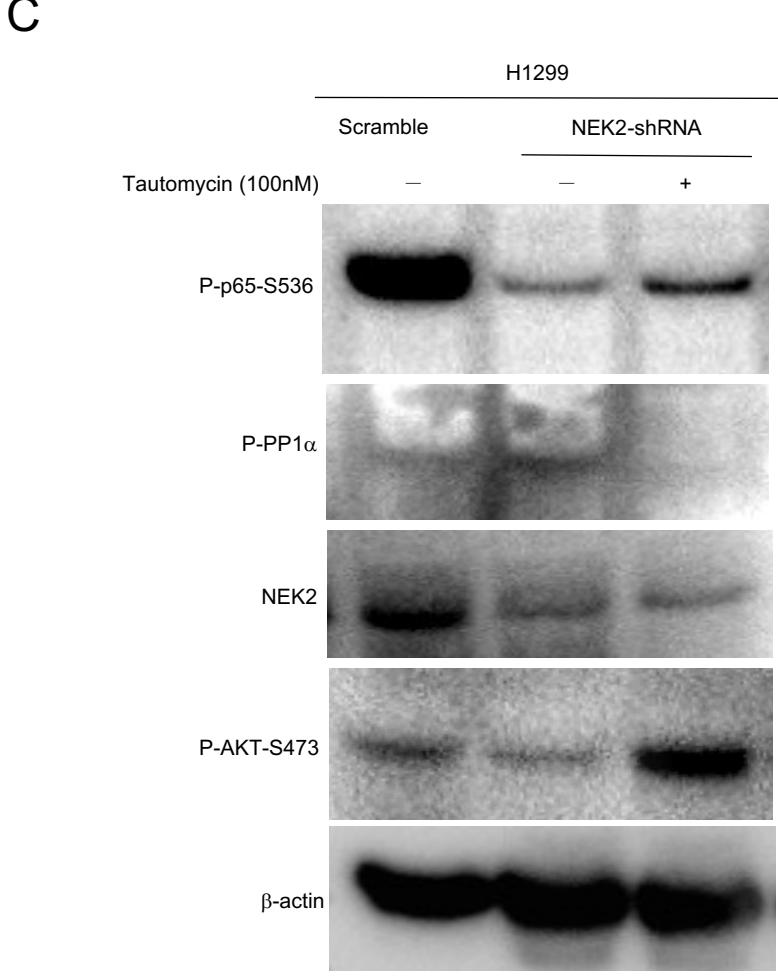
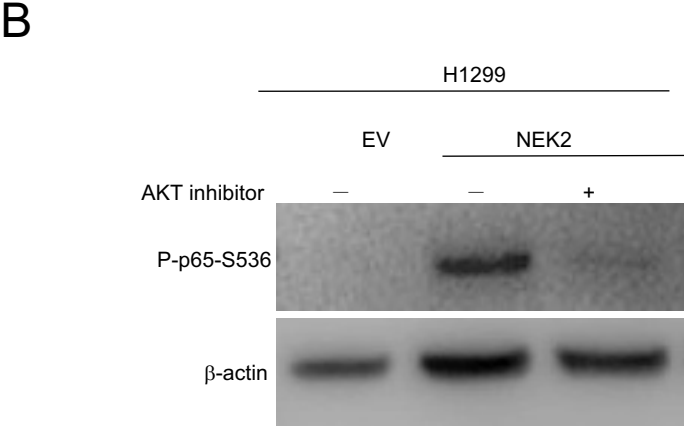
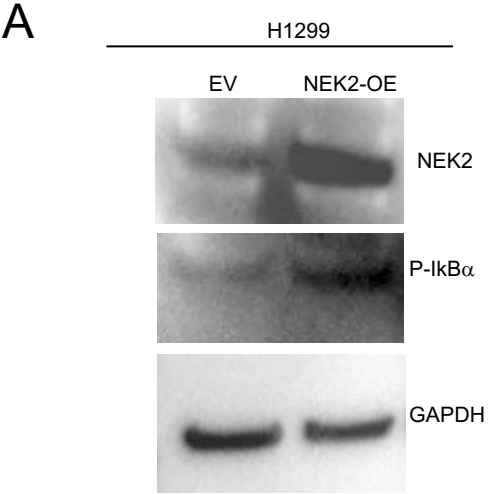
Supplemental Figure 2



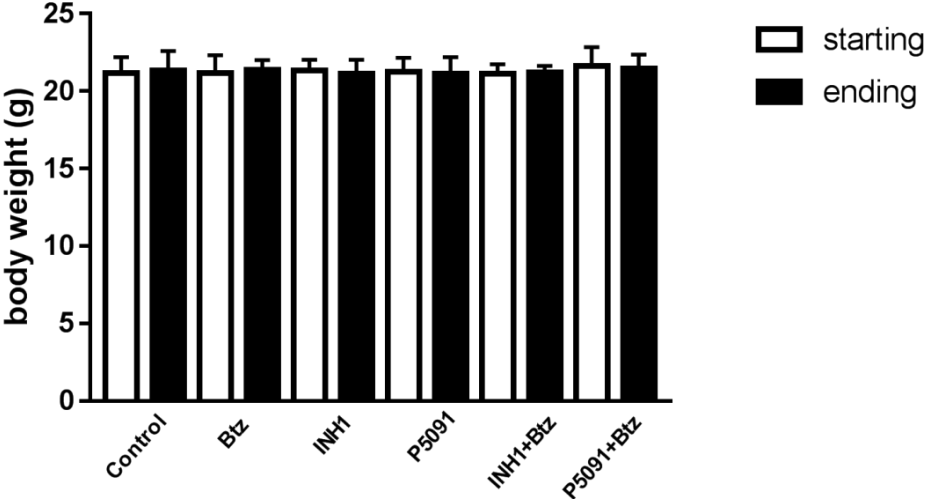
Supplemental Figure 3



Supplemental Figure 4



Supplemental Figure 5



Supplemental Table 2

Gene Symbol	p65 binding site*
<i>HPSE</i>	chr4:84,262,139-84,262,508
<i>HMGB1</i>	chr13:31,038,167-31,038,595
<i>LCP1</i>	chr13:46,739,634-46,740,590
<i>DARS2</i>	chr1:173,793,487-173,794,076
<i>NEDD1</i>	chr12:97,285,732-97,286,132
<i>PRKDC</i>	chr8:48,790,935-48,791,348
<i>CDCA2</i>	chr8:25,316,433-25,316,836
<i>CDC2</i>	chr14:50,810,854-50,811,264
<i>TPX2</i>	chr20:30,311,226-30,312,035
<i>SMC4</i>	chr3:160,117,270-160,117,688
<i>PPP1R12A</i>	chr12:80,328,739-80,329,571
<i>MKI67</i>	chr10:129,924,754-129,925,158
<i>PLAC8</i>	chr4:84,031,300-84,031,974
<i>DYRK2</i>	chr12:68,042,402-68,042,777
<i>PTEN</i>	chr10:89,623,165-89,623,590
<i>NLK</i>	chr17:26,382,567-26,382,950
<i>MLL</i>	chr6:168,106,886-168,107,255
<i>ALCAM</i>	chr3:105,085,406-105,086,113
<i>NFE2L1</i>	chr17:46,126,327-46,126,777
<i>GLG1</i>	chr16:74,486,191-74,486,614
<i>TGOLN2</i>	chr2:85,545,632-85,546,001
<i>CYLD</i>	chr16:50,775,635-50,776,546
<i>ALDH2</i>	chr12:112,211,700-112,212,283
<i>DCP2</i>	chr5:112,312,074-112,312,787
<i>TMED4</i>	chr7:44,621,599-44,622,341
<i>RC3H2</i>	chr9:125,667,614-125,668,078
<i>EXT2</i>	chr11:44,117,125-44,117,528
<i>IL15</i>	chr4:142,556,807-142,557,225
<i>TMEM179B</i>	chr11:62,554,693-62,555,096
<i>TTC17</i>	chr11:43,380,226-43,380,814
<i>C14ORF100</i>	chr14:59,910,898-59,911,267

* Human Feb. 2009 (GRCh37/hg19 from UCSC Genome Browser) Assembly, Track Name: GM12878+TNFa RELA Cell line: GM12878 (B cell)