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## **Supplemental Data**

## UV as an Amplifier Rather

# Than Inducer of NF-κB Activity

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### Figure S1

(A) Quantitation of the gels shown in Figure 2F.  $i\kappa b \alpha^{-/2}$  cells transduced with wild-type IkB $\alpha$  (IkB $\alpha$ WT, on left) or with Ser32/36Ala IkB $\alpha$  (IkB $\alpha$ AA, on right) were left untreated (black bars), treated with 60 J/m<sup>2</sup> UVC for 8 hours (dark gray bars), or TNF for 10 minutes as a control (light gray bars), and whole cell extracts subject to immunoprecipitation with anti-p65 antibody and immunoblot for IkB $\alpha$ . Levels of IkB $\alpha$  were normalized to the respective p65 levels and the ratio for untreated lanes were normalized to 1.0.

(**B**) Quantitation of the gels shown in Figure 2G.  $nf\kappa b^{-/-}$  cells transduced with either I $\kappa$ B $\alpha$ WT or I $\kappa$ B $\alpha$ AA were treated with CHX or UV and whole cell extracts subject to immunoblot for I $\kappa$ B $\alpha$ .

(C) Immunoblot for I $\kappa$ B $\alpha$  of whole cell extracts from  $nfkb^{-/-}$  ( $rela^{-/-}nf\kappa b1^{-/-}crel^{-/-}$ ) treated with 40 J/m<sup>2</sup> UVC for the indicated times.



To simulate UV-induced translational inhibition (Figures 3, 4, and 7), all protein synthesis parameters in the model were multiplied by a translational multiplier that linearly decreased from time zero to one hour from the modifier 1.0 to 0.5 (to make it 50% inhibition of protein synthesis, data based on results in Figure 1C). After one hour, the protein synthesis rates were held over time at 50% inhibition.

For the dose response curves in Figures 1, 5, and 6 the same tool was applied but the translational multiplier varied depending on the degree of translational inhibition indicated on the curve (dashed lines).



(A) Quantitation of Western blot gels shown in Figure 4B of  $nf\kappa b^{-/-}$  cells transduced with either I $\kappa$ B $\alpha$ WT or I $\kappa$ B $\alpha$ 6m treated with CHX. Band intensities were normalized to the untreated lane for each cell line.

(**B**) Immunoblot and quatitation of steady-state  $I\kappa B\alpha$  levels in  $i\kappa b\alpha^{-/2}$  cells trasduced with either  $I\kappa B\alpha WT$  or  $I\kappa B\alpha 6m$ . Levels of  $I\kappa B\alpha$  were normalized to their respective actin levels.

(C) Computational simulation of steady-state total cellular I $\kappa$ B $\alpha$  levels. In the "6m" model, the IKK-independent degradation rate of free I $\kappa$ B $\alpha$  was decreased 2-fold to mimic the effect of the 6m I $\kappa$ B $\alpha$  mutant.

(**D**) Computational simulations of NF- $\kappa$ B activation by UV-induced translational inhibition (40-50% translational inhibition, described in Supplemental Figure 2B). For "6m (slow deg)", red line, the IKK-independent free I $\kappa$ B $\alpha$  degradation rate was decreased 2-fold. For "high synthesis", blue line, I $\kappa$ B $\alpha$  degradation rates were as in the wild-type model, but the I $\kappa$ B $\alpha$  translation rate (mRNA to protein) was increased 1.9-fold, a degree to which the total cellular I $\kappa$ B $\alpha$  steady-state levels match the levels in the 6m model (1.3-fold higher than in the wild-type model).



Graph showing the calculated equilibrium levels of  $I\kappa B\alpha$ ,  $I\kappa B\beta$ , and  $I\kappa B\epsilon$  in the wild-type and slow flux models.

The specific alterations to parameters in the slow flux model as compared to the wildtype model are as follows: The IKK-independent degradation rate for free I $\kappa$ B $\alpha$  was divided by 2000, and for I $\kappa$ B $\beta$  and I $\kappa$ B $\epsilon$  they were divided by 3000 (all to make the IKKindependent degradation rate for free I $\kappa$ B equal to the respective IKK-dependent rate for bound I $\kappa$ B). The I $\kappa$ B $\alpha$  translation rate for I $\kappa$ B $\alpha$  was multiplied by 0.0752, the for I $\kappa$ B $\beta$ it was multiplied by 0.0177, and for I $\kappa$ B $\epsilon$  it was multiplied by 0.0283.







(A) In vitro IKK kinase assay of cytoplasmic extracts from wild-type cells treated with 1 ng/mL LPS ("LPS<sup>10</sup>") or co-treated with 1 ng/mL LPS and 40 J/m<sup>2</sup> UVC ("LPS<sup>10</sup> + UV"). Immunoblot for IKK $\beta$  is shown as a IP-efficiency and loading control. Quantitation from this gel was used as an IKK input for the computational simulations in Figure 7A.

(**B**) EMSA of nuclear extracts from  $i\kappa b\alpha^{-/2}$  cells transduced with either wild-type IkB $\alpha$  (IkB $\alpha$ WT) or Ser32/36Ala IkB $\alpha$  (IkB $\alpha$ AA) treated with 1 ng/mL only, 40 J/m<sup>2</sup> UVC only, or co-treated with LPS and UVC for the indicated times.

(C) EMSA of nuclear extracts from mouse keratinocytes treated with 1 ng/mL LPS or cotreated with 1 ng/mL LPS and 40 J/m<sup>2</sup> UVC. A graph showing quantitation of the NF- $\kappa$ B fold-induction (y-axis) at the 6-hour time point for each treatment regime is shown to the left of the gel.

(**D**) EMSA of wild-type cells treated with either 1 ng/mL IL-1, 40 J/m<sup>2</sup> UVC, 1 ng/mL TNF, co-treated with 1 ng/mL IL-1 and 40 J/m<sup>2</sup> UVC, or co-treated with 1 ng/mL TNF and 40 J/m<sup>2</sup> UVC for the indicated times

(E) In vitro IKK kinase assay of cytoplsmic extracts of wild-type cells treated with 1 ng/mL IL-1 or co-treated with 1 ng/mL IL-1 and 40 J/m<sup>2</sup> UVC. Immunoblot for IKK $\beta$  is shown as a IP-efficiency and loading control.





(A) Combined low dose CHX and low dose LPS hyper activate NF- $\kappa$ B. Nuclear extracts from wild-type cells treated with 0.1 µg/mL CHX ("CHX<sup>lo</sup>"), 1 ng/mL LPS ("LPS<sup>lo</sup>"), or co-treated with CHX and LPS for the indicated times were subject to EMSA. (B) Synergistic NF- $\kappa$ B activation by CHX and LPS requires IKK-mediated phosphoylation of I $\kappa$ B $\alpha$ . Nuclear extracts from  $i\kappa b \alpha^{-/-}$  cells transduced with either wild-type or Ser32/36Ala (I $\kappa$ B $\alpha$ AA) treated with 0.1 µg/mL CHX ("CHX<sup>lo</sup>"), 1 ng/mL LPS ("LPS<sup>lo</sup>"), or co-treated with CHX and LPS for the indicated times were subject to EMSA.



ER stress induced-translational inhibition depletes the pool of free I $\kappa$ B $\alpha$ . Immunoblot of whole cell extracts for I $\kappa$ B $\alpha$  from  $nf\kappa b^{-/-}$  MEFs transduced with wild-type I $\kappa$ B $\alpha$  treated with 40 or 400 nM thapsigargin or 0.1 mM or 1 mM DTT for the indicated time.