# Sequential DNA damage-independent and -dependent activation of NF-κB by UV

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NF-kB activation in response to UV irradiation of HeLa cells or of primary human skin fibroblasts occurs with two overlapping kinetics but totally different mechanisms. Although both mechanisms involve induced dissociation of NF-kB from IkBa and degradation of IkBa, targeting for degradation and signaling are different. Early IKBa degradation at 30 min to ~6 h is not initiated by UV-induced DNA damage. It does not require IKB kinase (IKK), as shown by introduction of a dominant-negative kinase subunit, and does not depend on the presence of the phosphorylatable substrate, IκBα, carrying serines at positions 32 and 36. Induced IkBa degradation requires, however, intact N- (positions 1-36) and C-terminal (positions 277-287) sequences. IKB degradation and NF-KB activation at late time points, 15-20 h after UV irradiation, is mediated through DNA damage-induced cleavage of IL-1a precursor, release of IL-1a and autocrine/ paracrine action of IL-1a. Late-induced IkBa requires the presence of Ser32 and Ser36. The late mechanism indicates the existence of signal transfer from photoproducts in the nucleus to the cytoplasm. The release of the 'alarmone' IL-1a may account for some of the systemic effects of sunlight exposure.

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# Introduction

In most types of cells, the dimeric transcription factor NF- $\kappa$ B, whose subunits belong to the Rel family, is kept in the cytoplasm through interaction with IkB proteins (reviewed by Bäuerle and Henkel, 1994; Bäuerle and Baltimore, 1996; Barnes and Karin, 1997). In response to one of numerous stimuli, e.g. cytokines, intracellular parasitic infections or phorbol ester, the inhibitory proteins (shown for I $\kappa$ B $\alpha$  and  $\beta$ ) are rapidly degraded, and NF- $\kappa B$  is released from its inhibitory association and taken up by the nucleus where it activates the transcription of, for example, genes important for the inflammatory and immune responses (reviewed by Verma et al., 1995; Bäuerle and Baltimore, 1996; Gilmore et al., 1996; Wulczyn et al., 1996; Barnes and Karin, 1997). Upon treatment with tumor necrosis factor  $\alpha$  (TNF $\alpha$ ), interleukin-1 $\alpha$  (IL-1 $\alpha$ ), or phorbol ester, I $\kappa$ B $\alpha$  is targeted for degradation by phosphorylation of the N-terminal serines at positions 32 and 36 (Brown *et al.*, 1995; Traencker *et al.*, 1995; DiDonato *et al.*, 1996). Recently an IkB kinase (IKK, CHUK) has been characterized and shown, by antisense deprivation and dominant-negative mutation, to be required for the activation of NF- $\kappa$ B by TNF $\alpha$ , IL-1 $\alpha$ , okadaic acid or phorbol ester (DiDonato *et al.*, 1997; Régnier *et al.*, 1997; Zandi *et al.*, 1997). The pathway to NF- $\kappa$ B may comprise still other protein kinases, e.g. NIK (Malinin *et al.*, 1997; Régnier *et al.*, 1997), presumably acting upstream of IKK. An alternative mechanism of NF- $\kappa$ B activation seems to occur in oxygen-deprived cells upon reoxygenation; IkB $\alpha$  appears to dissociate from NF- $\kappa$ B without degradation as a result of tyrosine phosphorylation (Imbert *et al.*, 1996).

Ultraviolet (UV) irradiation, UV-B or UV-C, of mammalian cells causes the activation of several transcription factors including NF-KB and the subsequent transcription of many genes, collectively called the UV response (reviewed by Bender et al., 1997; Herrlich et al., 1997). The original evidence for NF-κB regulation by UV stems from the observation that UV exposure of cells in culture or of mice induces transcription from the long terminal repeat (LTR) promoter of human immunodeficiency virus 1 (HIV-1; Valerie et al., 1988; Stein et al., 1989a; Morrey et al., 1991). The transcriptional activation of HIV-1 depends on the two NF-kB-binding sites located between positions -105 and -79 of the LTR, since their destruction abolishes the UV response. Also, NF-KB-binding sites cloned in front of a heterologous UV-non-responsive reporter render the construct UV-inducible (Stein et al., 1989a) and nuclear extracts isolated from UV-irradiated HeLa cells contain elevated NF-KB DNA-binding activity with maximal levels at 4 h after irradiation (Stein et al., 1989a). These experiments indicate that UV, like  $TNF\alpha$ or IL-1 $\alpha$ , targets NF- $\kappa$ B to the nucleus.

Our laboratory has been concerned with the signal transduction pathways elicited by UV, ultimately activating transcription factors (Schorpp et al., 1984; Herrlich and Karin, 1988; Karin and Herrlich, 1989; Mai et al., 1989; Sachsenmaier et al., 1994; Knebel et al., 1996). The analysis led from UV-responsive elements and transcription factors to signal transduction components and to the 'origins' of signaling. UV acts on more than one primary target molecule linked to signal transduction. Major primary target molecules generating UV-induced signaling are located at the plasma membrane (reviewed in Herrlich et al., 1997) leading to the ligand-independent activation of several receptor tyrosine kinases within fractions of a minute after UV irradiation, including receptors which may be relevant for the stimulation of NF-KB: the receptors for IL-1 $\alpha$  and TNF $\alpha$  (Sachsenmaier *et al.*, 1994; Rosette and Karin, 1996; Tobin et al., 1998). In agreement with UV induction through components in the plasma membrane, UV irradiation causes NF- $\kappa$ B activation in cytoplasts (Dévary *et al.*, 1993) and activates receptor tyrosine kinases, possibly also those relevant for NF- $\kappa$ B activation, in solubilized plasma membranes (Knebel *et al.*, 1996) or in cytosolic extracts supplied with cellular membranes (Simon *et al.*, 1994). UV-induced activation of receptor tyrosine kinases can be explained by transient or permanent inhibition of tyrosine phosphatases (Knebel *et al.*, 1996; S.Groß, A.Knebel, T.Tenev, A.Neininger, A.Deck, P.Herrlich, M.Gaestel and F.D.Böhmer, unpublished) which shifts the balance between negative and positive control elements towards increased signal flow.

Inconsistent with the notion of a signaling pathway originating in the plasma membrane is a series of experiments which can only be interpreted as involving the nucleus in signaling to NF-kB. For instance, far lower doses of UV-C (2 J/m<sup>2</sup>) were required for NF-κB-dependent HIV-LTR activation in cells from patients with Xeroderma pigmentosum group A (XPA) than in cells from healthy human individuals (20 J/m<sup>2</sup>; Stein et al., 1989a). Since these two types of cells differ only in their ability to repair UV-induced DNA photoproducts, the density of such lesions in the DNA must be a determinant of HIV transcription and presumably of NF-kB activation. Also in support of a role for DNA damage in NF-KB activation, lipofection of UV-irradiated pSV2-gpt DNA, but not of non-irradiated plasmid, into XPA cells stably transfected with a HIV-LTR-CAT plasmid induced CAT expression, which in turn was inhibited upon introduction by liposomes of the pyrimidine dimer excision enzyme, T4 endonuclease V (Yarosh et al., 1993).

Our present study aims at defining the mechanism of UV-induced activation of NF- $\kappa$ B and of I $\kappa$ B degradation in primary human skin fibroblasts and in HeLa cells. We show that UV causes I $\kappa$ B degradation by two totally different sequentially occurring mechanisms: an early phase does not involve the phosphorylation of I $\kappa$ B by I $\kappa$ B kinase while a late mechanism established in cultured cells beyond 15 h, requires the phosphorylatable I $\kappa$ B $\alpha$  Ser32 and Ser36, the DNA damage-induced processing of presynthesized IL-1 $\alpha$  precursor protein and release of IL-1 $\alpha$ . The action of secreted IL-1 $\alpha$  on other cells including inflammatory cells could explain some of the systemic effects of sun exposure.

#### Results

### Time-course of UV-induced NF-kB activity

Despite the fact that UV seemingly activates the IL-1 $\alpha$  and TNF $\alpha$  receptors in HeLa cells (Sachsenmaier *et al.*, 1994; Rosette and Karin, 1996), the time-course of NF- $\kappa$ B activation by UV differed from that induced by IL-1 $\alpha$  or TNF $\alpha$ . While IL-1 $\alpha$ , TNF $\alpha$  or phorbol ester caused rapid (15 min) and transient (60 min) I $\kappa$ B $\alpha$  degradation (not shown) and NF- $\kappa$ B activation (Figure 1A), UV radiation induced a delayed (30–60 min; Figure 1A) and long-lasting (still elevated at 42 h; Figure 1B) NF- $\kappa$ B response and reduction of I $\kappa$ B $\alpha$  levels (measured by Western blot, not shown; see below) in HeLa cells. Also in primary human skin fibroblasts, the response to TNF $\alpha$  was rapid while it took 60 min for the first appearance of active NF- $\kappa$ B after UV (not shown).



**Fig. 1.** Time-course of NF-κB activation. (**A**) HeLa cells were not treated or treated with TNFα (10 ng/ml), IL-1α (10 ng/ml), TPA (60 ng/ml) or UV-C (100 J/m<sup>2</sup>). Nuclear extracts were prepared at the indicated time points and equal amounts of proteins (5 µg) were analysed for NF-κB activity by electrophoretic mobility shift assay (EMSA). (**B**) HeLa cells were irradiated with 60 J/m<sup>2</sup> UV-C and analysed at the indicated time points for NF-κB activity as described in (A). The position of the shift produced by the p50/p65 heterodimer is indicated.

# DNA damage-independent and -dependent induction of NF-*k*B activity

As the induction of HIV-1 promoter activity depended on DNA photoproducts (Stein et al., 1989a), we first attempted to show similar dependence for UV-induced NF-KB activity. We exploited the specific difference in the repair kinetics (but not generation) of photoproducts between primary cells from a healthy human individual and those from a patient with XPA. As for the HIV-1-CAT reporter plasmid, NF-KB activity should be inducible in XPA cells at a lower dose than in wild-type cells, provided that DNA photoproducts were indeed intermediates and the repair time until harvest for NF-KB determination was sufficient to establish, by repair in the healthy cells, a damage-density difference. In wild-type human fibroblasts, 6-4-photoproducts are repaired with a half-life of 60-90 min, pyrimidine dimers with a half-life of 4-6 h (Friedberg et al., 1995). To be certain that sufficient DNA lesion density was reached, we determined in parallel the stabilization of p53 which is known to depend on DNA damage (Yamaizumi and Sugano, 1994). p53 accumulation, clearly visible at 6 h, was maximally induced by 5-10 J/m<sup>2</sup> in XPA cells and by 30 J/m<sup>2</sup> in wild-type fibroblasts (Figure 2, upper two panels, lanes 1-7) reflecting the photoproduct density difference between XPA and wild-type cells reached by repair in wild-type cells over the 6 h time period. This dose difference becomes of course even more pronounced with repair time (Figure 2, upper two panels, lanes 8–14).

However, NF- $\kappa$ B activation at 6 h after UV irradiation in the same extracts occurred with identical UV dose in



**Fig. 2.** Two mechanisms of UV-C-induced NF-κB activation. Wildtype fibroblasts and XPA fibroblasts were irradiated with the indicated doses of UV-C. Nuclear extracts were prepared 6 and 24 h after irradiation. For detection of p53 equal amounts of nuclear extracts (35 μg) were analyzed by Western blotting with the human p53specific monoclonal antibody Ab-2 (upper two panels). The position of the p53 protein is indicated. n.s., non-specific signal. For detection of NF-κB activity equal amounts of protein (5 μg) were analyzed by EMSA (lower two panels). The heterodimeric p50/p55 and the homodimeric p50/p50 complexes are indicated.

XPA and wild-type cells, reaching maximal activity at ~30 J/m<sup>2</sup> (Figure 2, lower two panels, lanes 1–7; note that in primary fibroblasts p50 is in excess, but not in HeLa cells, see Figure 1). In contrast, NF-kB activity measured at later time points, e.g. at 24 h, was induced in XPA cells by 5  $J/m^2$  while the dose required in normal cells remained high, 60 J/m<sup>2</sup> (Figure 2, lower two panels, lanes 8-14). These results imply that NF-KB activity in fibroblasts was induced by UV through two different mechanisms, one responsible for early activation, the other for delayed induction. Only the latter mechanism involved DNA photoproducts as necessary intermediates. The timecourse of NF-KB activation (Figure 1) may thus result from the fusion of at least two induction kinetics. Subsequently we explored which signal transduction pathways to NF- $\kappa$ B were exploited by the two types of UV induction. The early mechanism was studied in HeLa cells and in primary human fibroblasts in parallel while the delayed mechanism was resolved from the early induction in primary fibroblasts from a patient with XPA.

# The early mechanism of UV-induced NF-xB activation

Suramin resistance. The possibility that UV-induced ligand-independent receptor tyrosine kinase activation (Sachsenmaier *et al.*, 1994) was involved, was tested by the non-specific surface receptor poison, suramin, which abolishes UV-induced receptor tyrosine kinase activation (Sachsenmaier *et al.*, 1994; Iordanov *et al.*, 1997a). The early activation of NF- $\kappa$ B by UV was resistant to suramin, in contrast to that by growth factor containing conditioned medium (not shown). Thus a suramin-sensitive receptor tyrosine kinase was not involved in early UV induction of NF- $\kappa$ B in HeLa cells or in fibroblasts.



**Fig. 3.** Pretreatment with IL-1α does not interfere with UV-C-induced NF-κB activation. HeLa cells were not treated (–) or treated with IL-1α (10 ng/ml) or UV-C (100 J/m<sup>2</sup>) for the indicated times. These pretreatments were followed by a second treatment with either IL-1α (10 ng/ml) or UV-C (100 J/m<sup>2</sup>) where indicated. NF-κB activity of equal amounts of nuclear extract (5 µg) was analyzed by EMSA (upper panel). The amount of IkBα proteins was determined in cytoplasmic extracts by Western blotting with the anti-IkBα antibody sc-371 (lower panel).

No cross-refractoriness with IL-1 $\alpha$  or TNF $\alpha$ . The suramin resistance suggested that the receptors for IL-1 $\alpha$  or TNF $\alpha$ , both of which are activated by UV in HeLa cells and keratinocytes (Sachsenmaier et al., 1994; Rosette and Karin, 1996; Tobin et al., 1998), were not involved in the UV-activation of NF- $\kappa$ B. Nevertheless, we explored their putative involvement in a second type of approach. As shown previously (see also Sachsenmaier et al., 1994), most induced pathways become transiently refractory to a second stimulation by the same agent. Cross-refractoriness occurs if two stimuli address the same limiting signaling component. IL-1 $\alpha$  induced transient NF- $\kappa$ B activity (15) min; Figure 3, lane 2). After 2 or 8 h, NF-κB activity has disappeared almost completely (Figure 3, lanes 3 and 5) which begs the question as to whether re-treatment with a second stimulus re-induces NF-KB. Restimulation with IL-1 $\alpha$  did not activate NF- $\kappa$ B (Figure 3, lane 4). Pretreatment with either IL-1 $\alpha$  or TNF $\alpha$ , however, could not prevent UV-dependent rapid degradation of IkBa and NFκB activation [Figure 3, compare lane 7 (UV-induced after IL-1 $\alpha$  pretreatment) with lane 6 (UV only); data for TNF $\alpha$ not shown]. Thus components of signaling downregulated upon cytokine treatment, are not required for UV-induced signaling but are required for reinduction by cytokine. Restimulation with TNF $\alpha$  after pretreatment with IL-1 $\alpha$ or the reverse did, however, induce a second response (not shown), indicating that the two growth factor pathways involve different limiting components, both of which are not involved in the UV response.

We conclude that degradation of  $I\kappa B\alpha$  and activation of NF- $\kappa B$  at early time points after UV irradiation did not occur, at least not predominantly, through DNA damage nor through activation of components of the IL-1 $\alpha$  or TNF $\alpha$  receptor-driven pathways.

The early activation of NF- $\kappa$ B does not require the phosphorylation of I $\kappa$ B $\alpha$  Ser32 and Ser36 and cannot be inhibited by dominant-negative I $\kappa$ B kinase. The data presented so far suggest that the mechanism by which UV causes I $\kappa$ B $\alpha$  degradation, differs from that utilized by IL-1 $\alpha$  or TNF $\alpha$ . IL-1 $\alpha$  and TNF $\alpha$  are known to cause phosphorylation of I $\kappa$ B $\alpha$  at Ser32 and Ser36 (Figure 5A) which then leads to degradation through a ubiquitin-dependent pathway (Brown *et al.*, 1995; Chen *et al.*,



Fig. 4. ΙκBα Ser32 and Ser36 and CHUKL are required for TNFαbut not for UV-induced degradation. (A) HeLa cells were nontransfected (mock) or transiently transfected with epitope-tagged derivatives of the indicated IkBa cDNA expression vectors (2 µg). The cells were subsequently stimulated with UV-C (100  $J/m^2$ ) or with human TNFa (10 ng/ml) and harvested after 6 h or 30 min, respectively. The amount of  $I\kappa B\alpha$  proteins was determined in cvtoplasmic extracts by Western blotting with the anti-IKBa antibody sc-371. The positions of the exogenous and endogenous IKBa-proteins are indicated. (B) HeLa cells were transiently transfected with cDNA expression vector encoding epitope-tagged I $\kappa$ B $\alpha$  (2 $\mu$ g) or with the parental vector (pCMV4; 2 µg). The cells were stimulated and analyzed as described in (A). (C) HeLa cells were transiently transfected with a cDNA expression vector (1 µg) for epitope-tagged wild-type (wt) I $\kappa$ B $\alpha$  together with a cDNA expression vector (5  $\mu$ g) for IKK-β/CHUKwt or for IKK-β/CHUK with a mutated kinase domain (CHUKL-KD). The cells were stimulated and analyzed as described in (A). (D) Quantitative evaluation of  $I\kappa B\alpha$  levels by densitometry.

1995; Traencker *et al.*, 1995; DiDonato *et al.*, 1996). To investigate whether Ser32 and Ser36 were required for the UV response, we transiently expressed in HeLa cells (Figure 4) or XPA fibroblasts (see Figure 7) FLAG-tagged human wild-type or mutant I $\kappa$ B $\alpha$  driven by the cytomegalovirus (CMV) promoter. The exogenous wild-

type protein (larger than endogenous  $I\kappa B\alpha$ ) was degraded with similar efficiency and kinetics as the endogenous protein upon treatment of cells with either IL-1 $\alpha$  (not shown), TNF $\alpha$  or UV (Figure 4A and B, middle panel). Deleting 36 N-terminal amino acids ( $I\kappa B\alpha\Delta N$ ) stabilized IkB $\alpha$  and made it non-responsive to any treatment including UV (Figure 4A). The N-terminus is apparently required for recognition by the proteasome pathway (e.g. comprising the lysines at positions 21 and 22 which serve as ubiquitination substrates; Alkalay et al., 1995; Chen et al., 1995; Scherer et al., 1995; Baldi et al., 1996). A double point mutant of I $\kappa$ B $\alpha$  with Ser32 and Ser36 replaced by alanines (S32/36A), was resistant to TNF $\alpha$ - or IL-1 $\alpha$ induced degradation (shown for TNF $\alpha$  in Figure 4B, right panel; see also Figure 7), in agreement with published results (Brockman et al., 1995; Brown et al., 1995; Traencker et al., 1995). IκBαS32/36A was, however, still degraded in response to UV (Figure 4B, right panel; see quantitation in Figure 4D; see also Figure 7). Consistent with this result, transient co-transfection of HIV-CAT reporter with an expression vector encoding FLAG-tagged IkB $\alpha$ S32/36A reduced the response to TNF $\alpha$  but not to UV while co-expression of  $I\kappa B\alpha$  wild-type affected both inductions (not shown). Thus, UV-induced IkBa degradation does not appear to require phosphorylation of Ser32 and Ser36, in contrast to that induced by the cytokines.

A consequence of this result is that the UV-dependent degradation of IkBa may not involve IkB kinase-driven phosphorylation of the serines at positions 32 and 36 (DiDonato et al., 1997; Zandi et al., 1997). IKK does, however, phosphorylate a C-terminal serine (DiDonato et al., 1997). To test whether IKK was needed for the UV response, we aimed at inactivating the enzyme. The active form of IkB kinase consists of at least two subunits, IKKa and IKK $\beta$ , coupled by a leucine zipper (Zandi *et al.*, 1997). Both share ATP-binding motifs and contribute to catalytic activity. If UV could target IkB for degradation by a mechanism distinct from that requiring phosphorylation, a dominant-negative mutant of IkB kinase should not interfere with the early UV response of NF-kB. In our transfection conditions, a D $\rightarrow$ N mutant of IKK $\beta$ (CHUKL-KD) which is unable to bind ATP, severely inhibited IkB degradation after treatment of cells with TNF $\alpha$  (Figures 4C and D; note that these are transient transfections. The determination of endogenous IkB protein in this case reflects the behavior in the bulk of cells. IkB levels are therefore not visibly affected by dominantnegative IKK $\beta$ ). The dominant-negative mutant caused the same degree of inhibition of TNF $\alpha$ -induced I $\kappa$ B $\alpha$ degradation as the S32/36A mutation of IkB. The UV response was, however, not inhibited (Figures 4C and D) indicating that IKK was not involved in UV-induced signal transduction in the cells examined here, despite the fact that the response to TNF $\alpha$  involved IKK. This result also eliminates the involvement of IKK-dependent C-terminal phosphorylation.

*C-terminal sequence requirement*. Because of the reported  $I\kappa B\alpha$  tyrosine (Tyr42) phosphorylation after either reoxygenation of Jurkat cells or treatment with the tyrosine phosphatase inhibitor, pervanadate (Imbert *et al.*, 1996), we explored the possibility that  $I\kappa B\alpha$  could be directly phosphorylated at tyrosines rather than at the serines, in



Fig. 5. Amino acids between  $I\kappa B\alpha$  positions 277 and 287 are required for UV-induced degradation. (A) IxBa mutants. The upper graph shows a scheme of the IκBα protein with S32 and Ser36, and Tyr42 which is phosphorylated upon reoxygenation of cells. The ankyrin repeats are indicated in gray, the PEST sequence is indicated in black. HA is tagged to the N-terminal end of the protein. The lower graph shows the amino acid sequence between positions 247 and 317 and the deletion mutants which were used in this study. (B) HeLa cells were transiently transfected with 2  $\mu$ g of the indicated gene constructs encoding the HA-I $\kappa$ B $\alpha$  fusion proteins (lanes 2–16) or with 2  $\mu$ g of the parental vector pCMV4 (mock; lane 1). The cells were subsequently stimulated with UV-C (100 J/m<sup>2</sup>, 6 h) or with human TNF $\alpha$  (10 ng/ml, 15 min). The amount of I $\kappa$ B $\alpha$  proteins encoded by the exogenous gene constructs was determined in cytoplasmic extracts by Western blotting with a monoclonal antibody specific for the HA epitope (clone 12 CA 5). (C) Ser283 is not required for UV- or TNFαinduced IkB $\alpha$  degradation. Ser283 was mutated to alanine, either in the complete I $\kappa$ B $\alpha$  protein (lanes 17–19) or in the deletion mutant HA-I $\kappa$ B $\alpha$  1–287 (lane 20–22). The mutated gene constructs were analyzed for IkBa expression as in (B). n.s., non-specific signal.

response to UV. In both HeLa and Jurkat cells treated with pervanadate,  $I\kappa B\alpha$  phosphorylated at tyrosines was detected with either an  $I\kappa B\alpha$ -specific or a tyrosine phosphate-specific antibody. After UV irradiation, however,  $I\kappa B\alpha$  was not phosphorylated at tyrosines either in Jurkat or in HeLa cells (data not shown).

To screen for other sequences in IkB $\alpha$  that might be relevant for UV-induced turnover, we focused on the C-terminal region which is, in addition to the N-terminus, required for the TNF $\alpha$ -induced degradation (Aoki *et al.*, 1996; Sun *et al.*, 1996). The C-terminus harbors a QLrich region (aa 264–277) and a PEST sequence (aa 281– 304) including several putative casein kinase II phosphorylation sites (Figure 5A) which seem not to exert a regulatory role (Barroga et al., 1995; Brown et al., 1995; Rodriguez et al., 1995; Whiteside et al., 1995; Aoki et al., 1996; Krappmann et al., 1996; Lin et al., 1996; McElhinny et al., 1996; Schwarz et al., 1996; Sun et al., 1996). In accordance with published observations,  $TNF\alpha$  could not trigger degradation of a long C-terminal truncation (I $\kappa$ B $\alpha$ aa 1–263, Figure 5B). Also UV did not induce degradation of this mutant protein. To dissect the area of the C-terminus absent in the 1-263 truncation, progressive deletions were tested. These deletion mutants revealed loss of UV-induced destabilization of I $\kappa$ B $\alpha$  prior to that by TNF $\alpha$  (Figure 5B) or IL-1 $\alpha$  (not shown). Clearly, deletion to position 277 had no effect on TNF $\alpha$ -induced turnover (Figure 5B) while all deletions reaching into the stretch between positions 277 and 287 (e.g.  $I\kappa B\alpha$  1–284; not shown) were stable in response to UV. IkBa 1-287 was, however, destabilized by UV. Since the stretch between positions 277 and 287 contains one serine at position 283, a putative target of casein kinase II (Barroga et al., 1995; Lin et al., 1996; McElhinny et al., 1996; Schwarz et al., 1996) and of a lipopolysaccharide-inducible protein kinase (Kuno et al., 1995), and since one might expect UV-induced phosphorylation to be the initiating event, we pointmutated this serine (S283A). In the context of the complete or truncated (to 287) IkBa protein, UV destabilized I $\kappa$ B $\alpha$ S283A (Figure 5C), suggesting, at least, that Ser283 does not play a prominent role in UV-induced turnover.

Thus we can conclude that the early onset of NF- $\kappa$ B activation after UV irradiation of primary human skin fibroblasts and HeLa cells does involve prior degradation of I $\kappa$ B, but by a mechanism different from that used by TNF $\alpha$  or IL-1 $\alpha$ . The degradation mechanism involves the N-terminal region, presumably the N-terminal lysines, and the stretch between amino acid positions 277 and 287. The C-terminal requirements also differ from those needed in TNF $\alpha$ -dependent degradation. Ser32, Ser36 and Ser283 as well as tyrosine phosphorylation are irrelevant for the early UV-induced turnover.

# The late mechanism of UV-induced NF-KB activation

DNA damage-mediated degradation of IKBa and activation of NF- $\kappa B$  are mediated through the release of IL- $1\alpha$ . The late induction of NF- $\kappa$ B activity can be dissociated from the early induction in XPA fibroblasts. In contrast to HeLa cells, human fibroblasts appear to carry an excess of p50 such that p65/p50 heterodimers and p50 homodimers are generated. Antibody supershifts are shown in Figure 6A; compare the constituents of NF-kB gelshifts of HeLa and XPA cells. Specificity of the gelshift and of the antibodies was ascertained by control gelshifts using an Oct-1 oligonucleotide and by adding an excess of specific or non-related oligonucleotides (Figure 6A). Induced NF-KB activity in XPA fibroblasts was resolved into two bands which, according to supershift experiments, represent p65/p50 and p50/p50 (Figure 6A). Also, the activity induced in HeLa cells was resolved into two complexes, which according to the supershifts can be identified as a major p65/p50 heterodimer and a minor p50/ p50 homodimer. It is not clear why the HeLa homodimer migrated slower than that of XPA cells. UV enhanced only the heterodimer in HeLa cells (Figures 1 and 6A),



Fig. 6. Composition of NF-KB, kinetics and IL-1a-dependence of the late UV response of NF-KB. (A) In order to investigate the composition of the complexes obtained, extracts of HeLa cells irradiated with 40 J/m<sup>2</sup> and harvested at 6 h, and extracts from XPA cells irradiated with 5 J/m<sup>2</sup> and harvested at 24 h, were preincubated with the p50-specific antibody sc-1190X or with the p65-specific antibody sc-372. To determine specificity of binding, we used a 30-fold excess of the non-labeled NF-kB-binding oligonucleotide or of the Oct-1-binding oligonucleotide as competitors. (B) Late activation of NF-KB in XPA cells with low UV doses. Wild-type and XPA fibroblasts were irradiated with 30 and 5 J/m<sup>2</sup>, respectively. At the indicated time points nuclear extracts were prepared and equal amounts of protein (5 µg) were analyzed for NF-KB activity by EMSA. The heterodimeric p50/p65 and the homodimeric p50/p50 complexes are indicated. (C) IL-1 $\alpha$  mediates the late UV signal to NF-KB. XPA fibroblasts were not irradiated or irradiated with UV-C as indicated. Immediately after irradiation the cells were treated with 10 µg/ml of neutralizing antibodies directed against TNFa (aTNFa), IL-1 $\alpha$  ( $\alpha$ IL-1 $\alpha$ ) or both ( $\Sigma$ Ab). As controls the cells were treated with an unrelated immunoglobulin ( $\alpha$ IgG) or not treated with antibodies. Six h (30 J/m<sup>2</sup>) and 24 h (5 J/m<sup>2</sup>) after irradiation nuclear extracts were prepared and equal amounts of protein (5 µg) were analyzed for NF-KB activity by EMSA.

but hetero- and homodimers in fibroblasts (Figures 2 and 6).

With 5 J/m<sup>2</sup> an early induction in XPA cells was below detection level, while the DNA damage-dependent activation occurred with a lag period of 20 h (Figure 6B). With 30 J/m<sup>2</sup> in wild-type fibroblasts, one may expect both mechanisms to occur. However, the early DNA damage-independent response obviously concealed, by merging, the second wave of induction (Figure 6B). What causes the delay in XPA cells irradiated with 5 J/m<sup>2</sup>



**Fig. 7.** IκBα Ser32 and Ser36 are required for the late UV-induced IκBα degradation. XPA fibroblasts were transiently transfected with a IκBα cDNA expression vector (2 µg) encoding the FLAG-tagged wild-type IκBα protein or the Ser32/36Ala mutant of IκBα. The cells were mock treated (lanes 1, 6, 8 and 13) or treated with TNFα (10 ng/ml), IL-1α (10 ng/ml), or UV-C (100 or 10 J/m<sup>2</sup>). Six or 24 h after irradiation and 30 min after treatment with TNFα or IL-1α cytoplasmic extracts were prepared and equal amounts of proteins (80 µg) were analyzed for the amounts of exogenous and endogenous IκBα proteins by Western blotting with the anti-IκBα antibody sc-371. The positions of the exogenous and endogenous IκBα-proteins are indicated.

of UV? After all, DNA damage by UV is inflicted instantaneously. An interesting possibility has been suggested by previous experiments showing UV-induced and apparently DNA damage-dependent synthesis and release of cytokines (termed EPIF, Schorpp et al., 1984; or UVIS, Rotem et al., 1987; damage dependence, Yarosh et al., 1993; M.Litfin and H.J.Rahmsdorf, unpublished). UVirradiated HeLa cells for instance release IL-1 $\alpha$  and bFGF (Krämer et al., 1993). UV-induced release, accumulation in the medium and autocrine or paracrine action of cytokines could well account for the slow time-course of NF-KB activation. To test this possibility, we examined whether we could interfere with the induction of NF- $\kappa B$ in XPA cells by neutralizing anti-cytokine antibodies. Of several antibodies tested, only those directed against IL- $1\alpha$  could obliterate the delayed response to 5 J/m<sup>2</sup> UV in XPA fibroblasts (of both hetero- and homodimers, Figure 6C, lanes 10 and 11; controls shown with antibodies to TNF $\alpha$ , lane 9, and IgG, lane 8). The early induction by 30 J/m<sup>2</sup> was not affected (Figure 6C, compare lanes 5 and 6). Thus, these data indicate that IL-1 $\alpha$  release is an intermediate in the delayed DNA damage-dependent UV response and that IL-1 $\alpha$  is the predominant cytokine secreted from UV-irradiated primary fibroblasts, mediating NF-KB activation.

Requirement for  $I\kappa B\alpha$  Ser32 and Ser36. If IL-1 $\alpha$  mediated the late DNA damage-dependent UV signal to NF- $\kappa$ B, late I $\kappa$ B $\alpha$  degradation should depend on the induced phosphorylation at Ser32 and Ser36. Introduced transiently into XPA fibroblasts, FLAG-tagged I $\kappa$ B $\alpha$ S32/36A was resistant to TNF $\alpha$ , IL-1 $\alpha$  and to UV determined at 24 h, compatible with its being mediated by IL-1 $\alpha$ , while at an early time point (6 h) it was degraded in response to UV (Figure 7; compare mutant lanes with wild-type lanes: e.g. 14 with 7, 9 with 2 etc.; compare also with the behavior of the endogenous I $\kappa$ B $\alpha$  which migrates faster than FLAG-tagged I $\kappa$ B $\alpha$ ; the data correspond to those obtained in HeLa cells in Figure 4).

IL-1 $\alpha$  precursor cleavage in the cytoplasm. The question remains as to how DNA damage can induce IL-1 $\alpha$  release. We have reported earlier that IL-1 $\alpha$  transcription is stimulated in HeLa cells upon UV irradiation (Krämer *et al.*, 1993). In XPA cells an effect on transcription was

barely detectable (not shown). Nevertheless, IL-1 $\alpha$  protein was released as revealed by immunoprecipitation with IL-1 $\alpha$ -specific antibodies (not shown). Since IL-1 $\alpha$  is synthesized as a precursor protein, we asked whether UVinduced DNA damage enhanced cleavage of the precursor. To improve the detection of IL-1 $\alpha$  precursor, we transiently transfected into XPA fibroblasts an expression clone encoding the FLAG-tagged IL-1a precursor protein. UV irradiation caused significant precursor cleavage as determined by Western blot experiments (Figure 8A). UV irradiation caused the decrease of the exogenous precursor protein coinciding with an increase in tagged IL-1 $\alpha$  followed by cytokine release into the medium. The cleavage occurred with an amazingly long lag period of 18 h, suggesting that the delay in NF-KB activation was caused by steps preceding IL-1 $\alpha$  processing rather than IL-1 $\alpha$  accumulation in the medium.

In an attempt to define the preceding steps we used a number of inhibitors. As one might have expected from the long lag period, UV-induced IL-1 $\alpha$  processing was sensitive to cycloheximide (not shown), indicating the need for protein synthesis. The final step in the processing of IL-1 $\alpha$  precursor involves proteolytic cleavage. The cysteine protease calpain has been proposed to be the processing enzyme (Kobayashi et al., 1990; Carruth et al., 1991; Kavita and Mizel, 1995). Inhibitors of this class of proteases, E64-d, Z-D.dcbmk and Z-VAD.fmk (Mehdi, 1991; Zhu et al., 1995; Nicholson and Thornberry, 1997), inhibited the cleavage and release of IL-1 $\alpha$  (shown for Z-VAD.fmk in Figure 8B). In keeping with the essential role of IL-1 $\alpha$  in the delayed UV response, the inhibitors also blocked the delayed UV-induced activation of NF- $\kappa B$  (Figure 8C, compare lanes 10–12 with lane 9). These inhibitors are likely to interfere with IL-1 $\alpha$  processing alone rather than with p105 cleavage, as E64-d has been shown not to affect p105 (Palombella et al., 1994).

Is the release of the alarmone IL-1 $\alpha$  by enhanced processing the consequence of apoptosis or an independent process? The time-course suggests a late event. IL-1 $\alpha$  processing preceded phosphatidylserine presentation on the cell surface (annexin V staining, not shown) which is an early sign of apoptosis. Only 5% of the cells showed annexin V staining at 24 h. This number of cells could not account for the degree of cleavage of IL-1 $\alpha$  precursor in the transfectants. Also the magnitude of the NF- $\kappa$ B response in XPA cells at 24 h after UV suggests that a considerable part of the response occurs in cells by autocrine stimulation either prior to apoptosis or in surviving cells.

### Discussion

Here we have shown that UV irradiation of primary human skin fibroblasts and of HeLa cells in culture activates the transcription factor NF- $\kappa$ B through two different pathways, both of which lead to I $\kappa$ B degradation. Early activation within the first 6 h after UV does not depend, at least mainly, on UV-induced DNA photoproducts and does not utilize components known to participate in cytokinedependent NF- $\kappa$ B activation such as IKK and its I $\kappa$ B substrate Ser32 and Ser36. Sequences in the N- and C-termini of I $\kappa$ B, however, need to be intact. Relevant UV absorption by a yet unknown suramin-resistant mem-



Fig. 8. The UV-induced release of IL-1 $\alpha$  depends on the cleavage of IL-1 $\alpha$  precursors. (A) XPA fibroblasts were transiently transfected with a cDNA expression vector (5  $\mu$ g) encoding the FLAG-IL-1 $\alpha$ precursor fusion protein. Twenty-four hours after transfection the cells were not irradiated or irradiated with 5 J/m<sup>2</sup>. At the indicated time points cell lysates were prepared and immunoprecipitated with anti-IL- $1\alpha$  antiserum (#1190) followed by Western blot with an anti-IL-1 $\alpha$ polyclonal antibody (AB-200NA). The two IL-1α precursor proteins are indicated. The slow-migrating band represents the Ig heavy chain, the n.s. (non-specific signal) band served as loading control. (B) XPA fibroblasts were transiently transfected with an IL-1\alpha-precursor expression vector as in (A). Immediately after irradiation the cells were not treated or treated with the caspase inhibitor Z-VAD.fmk (50 μM). Twenty-four hours later mature IL-1α was immunoprecipitated from the culture medium using a FLAG epitope-specific antibody (sc-807; Santa Cruz). The level of IL-1a detected in the culture medium of non-treated cells was set at 100%. (C) Several caspase inhibitors prevent late NF-KB activation. XPA fibroblasts were not treated, treated with IL-1 $\alpha$  (10 ng/ml) or irradiated with 5 J/m<sup>2</sup> or 30 J/m<sup>2</sup> UV. Caspase inhibitors Z-VAD.fmk (50 µM) or Z-D.dcbmk (100  $\mu$ M) and the calpain inhibitor E 64-d (50  $\mu$ M) were added to the cells immediately after treatment as indicated. Nuclear extracts were prepared 30 min after IL-1a addition, 6 h after irradiation with 30 J/m<sup>2</sup> and 24 h after irradiation with 5 J/m<sup>2</sup> and analyzed for NF- $\kappa$ B activity by EMSA. The positions of the shifts produced by p50/p65 heterodimers and p50/p50 homodimers are indicated.

branal or cytoplasmic chromophore may link to a signal transduction component that reaches  $I\kappa B$  (see also below). In contrast, the delayed UV response of NF- $\kappa B$  requires DNA damage as an intermediate which sets off a signaling chain out of the nucleus leading to the processing and

release of the alarmone IL-1 $\alpha$ , followed by autocrine/ paracrine induction of the IL-1 $\alpha$  signaling pathway to NF- $\kappa$ B. This late mechanism of induction is measured predominantly in the UV-induced expression of NF- $\kappa$ B reporter with long-term accumulation of CAT (Stein *et al.*, 1989a; Yarosh *et al.*, 1993). Similarly, late activation of other transcription factors mediating UV-induced expression of urokinase, metallothionein or collagenase (Miskin and Reich, 1980; Schorpp *et al.*, 1984; Stein *et al.*, 1989a,b) may be explained by IL-1 $\alpha$  release. The dual mechanism of NF- $\kappa$ B activation thus solves the apparent discrepancy between previous studies proposing nuclear and non-nuclear pathways (Stein *et al.*, 1989a; Devary *et al.*, 1993).

#### The early pathway to NF-κB

Following irradiation of HeLa cells (Figure 1) or human primary fibroblasts (Figure 6B) with UV, NF- $\kappa$ B activity is detectable with a lag period of 30–60 min, followed by massive increases at 2–6 h. This time-course is slow in comparison with the activation of other early UV targets (Devary *et al.*, 1992; Hibi *et al.*, 1993; Radler-Pohl *et al.*, 1993; Sachsenmaier *et al.*, 1994; Gupta *et al.*, 1995; Livingstone *et al.*, 1995; Raingeaud *et al.*, 1995; van Dam *et al.*, 1995; Price *et al.*, 1996; Iordanov *et al.*, 1997a). It is also slow in comparison with NF- $\kappa$ B activation after TNF $\alpha$  or IL-1 $\alpha$ . The delay of NF- $\kappa$ B activation (30–60 min) resembles the induction of signaling through collagen receptors (Vogel *et al.*, 1997). It is not known which ratelimiting step causes the delay.

The time-course difference (between TNF $\alpha$  and UV induction of NF- $\kappa$ B) matches a difference in the mechanism. I $\kappa$ B $\alpha$  is phosphorylated at Ser32 and Ser36 upon TNF $\alpha$  treatment, but not after UV. Nevertheless it is probably ubiquitinated since the N-terminus (with the lysine residues) is required for degradation. We do not yet know how ubiquitination is triggered by UV. Different sequence requirements in the I $\kappa$ B $\alpha$  C-terminus between TNF $\alpha$  and UV suggest recognition by a different ubiquitin-conjugating enzyme. Interestingly, the appearance of p50 homodimers is also enhanced by UV irradiation, demanding that the UV-induced early mechanism encompasses p105 processing.

The level of IkB is the net result of synthesis and degradation. Since  $I\kappa B\alpha$ , whether complexed with NFκB (Krappmann et al., 1996; van Antwerp and Verma, 1996) or in excess of NF-KB, is an intrinsically unstable protein (IkBa mRNA half-life was 30-45 min, protein half-life was 120-160 min for endogenous, complexed I $\kappa$ B, and 45 min for overexpressed, free I $\kappa$ B; not shown), a block of resynthesis would cause disappearance of  $I\kappa B\alpha$ and a delayed release and activation of NF-kB. UV treatment of cells indeed affects both transcription and translation (Sauerbier and Hercules, 1978; Iordanov et al., 1998). One could hypothesize that this inhibition was the cause of a net loss of IkB. A simply passive induction of NF- $\kappa$ B by loss of I $\kappa$ B $\alpha$  due to UV-induced translational inhibition is however highly unlikely: (i) even  $100 \text{ J/m}^2$ of UV inhibited overall translation by only 50% (not shown, and Iordanov et al., 1998); (ii) inhibition of IKB translation to a similar degree by emetine, which is not supposed to cause ribosome-borne signal transduction (Iordanov et al., 1997b), could not mimic NF-KB induction by UV (not shown); (iii) a non-specific reduction of  $I\kappa B$ by blocking synthesis should not be vulnerable to the introduction of  $I\kappa B$  mutations. However, several mutant proteins are stable after UV irradiation, e.g.  $I\kappa B\alpha\Delta N$ and  $I\kappa B\alpha 1-277$ , whereas slightly different mutants, e.g.  $I\kappa B\alpha 1-287$ , expressed at similar levels, are readily degraded upon UV irradiation (Figure 5B). Moreover, all mutant proteins were degraded with similar kinetics in non-irradiated cells treated with the protein synthesis inhibitors, cycloheximide or emetine (not shown).

Translational inhibition by UV-C is also not sufficient to explain the late induction of NF- $\kappa$ B-dependent promoters which were measured by CAT expression (Stein *et al.*, 1989a). Further, translational inhibition should not depend on DNA damage-repair; the dose of UV required should be identical between wild-type and XPA cells. However, 2 J/m<sup>2</sup> UV-C induce a maximum of HIV-1 transcription in XPA cells while 30 J/m<sup>2</sup> are required in wild-type cells.

Similarly, loss of I $\kappa$ B by transcriptional inhibition cannot explain UV-induced NF- $\kappa$ B activation; there is no major drop of I $\kappa$ B $\alpha$  mRNA in HeLa cells after irradiation even with doses as high as 100 J/m<sup>2</sup> (not shown), at which dose I $\kappa$ B $\alpha$  protein is degraded and NF- $\kappa$ B is activated. In summary, these data suggest the existence of a specific early signaling pathway which destabilizes I $\kappa$ B $\alpha$  and which is distinct from that originating from known surface receptors stimulated by their ligands.

It is puzzling that UV irradiation induces activation of the receptors for TNF $\alpha$  and IL-1 $\alpha$  in HeLa cells and in mouse keratinocytes (Sachsenmeier et al., 1994; Rosette and Karin, 1996; Tobin et al., 1998). Nevertheless the pathways to UV-induced early induction of NF-κB differ from those elicited by the ligands. This conclusion is based on the finding that IKK and the substrate serines are not required and that there is no cross-refractoriness of pathways. Differences in signal transduction between different types of cells, as have been found for UVinduced AP-1 activation (Sachsenmaier et al., 1994; Huang et al., 1997), are possible but do not seem to apply here, as keratinocytes, where TNF $\alpha$  and IL-1 $\alpha$  receptor activations have been measured, and HeLa cells should share pathways. The pathways triggered by the  $TNF\alpha$ receptor are not totally clear. While TNFa induces JNK activity through receptor-TRADD/TRAF-2, NF-KB seems to be activated mainly by a different, as yet unknown, pathway (Lee et al., 1997; Yeh et al., 1997). The report that UV-B induction of NF-kB was inhibitable by dominantnegative TNFα-receptor and TRAF-2 (Tobin et al., 1998) could indicate that UV-B activates the receptor and that this activation is required for UV induction along with a second signal, not involving IKK. A role for the  $TNF\alpha$ receptor even in the UV-induced JNK activation has also been suggested recently (Iordanov et al., 1998), since inhibitors of the generation of ribosome-borne signaling blocked JNK activation. Thus, a more complete understanding of TNFa receptor activation by UV compared with ligand activation is required to dissect the pathways to NF-KB.

Interestingly, suramin-resistant activation of cellular signal transduction has also been observed in the UV-induced activation of p38/HOG1 (Iordanov *et al.*, 1997a), in contrast to the suramin-sensitive activation of Erk in the same cells. NF- $\kappa$ B is, however, not downstream of

p38 since a p38 inhibitor, SB203580 (Cuenda *et al.*, 1995), could not block UV-induced  $I\kappa B\alpha$  degradation (not shown).

### The delayed pathway to NF-*k*B

In contrast to the transient activation of NF- $\kappa$ B by most inducers, activation by UV irradiation is prolonged over 24–48 h. As shown by dose differences between wildtype and XPA cells, the expression of an NF- $\kappa$ B-dependent reporter determined at 42 h after UV irradiation (Stein *et al.*, 1989a) and the delayed I $\kappa$ B $\alpha$  degradation and NF- $\kappa$ B activation described here, depend on UV-induced DNA damage.

Work on UV-induced transcription of collagenase and metallothionein, as well as on UV-induced stabilization of p53, has identified DNA damage in the transcribed strand of active genes as relevant for the signaling process (Yamazumi and Sugano, 1994; Blattner *et al.*, 1998) and thus suggested that stalled transcription complexes may generate the signal. Subsequently a chain of events is induced that ultimately, after a considerable lag period, activate proteases, one of which processes the IL-1 $\alpha$  precursor. Mature IL-1 $\alpha$  then activates, through its membrane receptor, IKK to phosphorylate I $\kappa$ B $\alpha$  and to initiate degradation.

The nature of the signal generated by stalled transcription complexes is unfortunately not yet known. We do not wish to speculate as to its nature as this paper does not contribute to the question of signal generation except for indicating the existence of a signal flow out of the nucleus into the cytoplasm. Whereas DNA damage by radiation is inflicted instantaneously, IL-1 $\alpha$  processing occurs after a long lag period. The susceptibility to protease inhibitors could mean that the processing itself was inhibited, or that IL-1 $\alpha$  was downstream of caspase action induced as part of the apoptotic program. Interestingly CD95-induced JNK activation was also sensitive to inhibition of cysteine proteases (Cahill *et al.*, 1996).

The delayed release of an alarmone, IL-1 $\alpha$ , may account for several delayed responses after UV irradiation: transcription of collagenase, metallothionein and, in part, mutagenesis (Maher *et al.*, 1988; Boesen *et al.*, 1992; Krämer *et al.*, 1993). Conditioned medium from UVirradiated cells induces the transcription of plasminogen activator, collagenase I, HIV-1 and metallothionein IIA (Schorpp *et al.*, 1984; Rotem *et al.*, 1987; Stein *et al.*, 1989b; Krämer *et al.*, 1993; Yarosh *et al.*, 1993). IL-1 $\alpha$ release is certainly a good candidate for some of the systemic effects of sun exposure. Both precursor cleavage and free IL-1 $\alpha$  could serve as targets for strategies that would prevent unwanted side-effects of radiation and of treatments with other DNA damaging agents, for example in the course of cancer therapy.

# Materials and methods

### Plasmids

The I $\kappa$ B $\alpha$  expression plasmids pCMV-FLAG-I $\kappa$ B $\alpha$  wt and pCMV-FLAG-I $\kappa$ B $\alpha$  Ser32/36Ala, in which Ser32 and Ser36 were mutated to alanines (Scherer *et al.*, 1995), were obtained from Dr Dean Ballard (Vanderbilt University, Nashville, TN). The I $\kappa$ B $\alpha$  C-terminal truncation mutants (Sun *et al.*, 1996) pCMV-HA-I $\kappa$ B $\alpha$  (1–317), pCMV-HA-I $\kappa$ B $\alpha$  (1–263), pCMV-HA-I $\kappa$ B $\alpha$  (1–277), pCMV-HA-I $\kappa$ B $\alpha$  (1–287) and pCMV-HA-I $\kappa$ B $\alpha$  (1–297) were generously given by Dr Warner Greene

(University of California, San Francisco, CA). Site-specific mutations of pCMV-HA-I $\kappa$ B $\alpha$  (1–317/Ser283Ala) and pCMV-HA-I $\kappa$ B $\alpha$  (1–287/Ser283Ala) and a D145N mutant of IKK $\beta$  (CHUKL) unable to bind ATP were created by PCR cloning and site-directed mutagenesis (Stratagene). The pRc/RSV-pIL1 $\alpha$ FLAG encoding the IL-1 $\alpha$  precursor (Siders *et al.*, 1993) was a gift of Dr S.Mizel (Wake Forest University, NC).

#### Cell culture and transfections

Cells were grown at 37°C and 6% CO<sub>2</sub>. HeLa tk<sup>-</sup> cells (Angel *et al.*, 1987) were maintained in Dulbecco's modified Eagle's medium (DMEM) supplemented with 8% fetal calf serum (FCS). Primary human fibroblasts from a healthy donor (a gift from Dr Eife, München) and fibroblasts from a patient with XPA (a gift of Dr J.Cleaver, CA) were grown in DMEM supplemented with 10 and 20% FCS, respectively. The human leukemic T cell line, Jurkat, was maintained in RPMI 1640 containing 10% heat-inactivated FCS, 1 mM sodium pyruvate and 2 mM L-glutamine (Gibco-BRL). All culture media were supplemented with 100 U/ml penicillin and 100  $\mu$ g/ml streptomycin (from Gibco-BRL).

HeLa cells were plated 20-24 h prior to transfection at a density of  $1 \times 10^6$  cells per 100 mm dish. Transfection was performed using DEAE-Dextran (Kawai and Nishizawa, 1984). Fibroblasts were plated 20-24 h prior to transfection at a density of  $8 \times 10^5$  cells per 100 mm dish. Transfection was performed using Lipofectamine<sup>TM</sup> Reagents (Gibco-BRL).

#### Antibodies, reagents and UV irradiation

The following antibodies were used: anti-IkBa rabbit polyclonal (sc-371, Santa Cruz); anti-phosphotyrosine rabbit monoclonal (PY-20, Transduction Labs); anti-FLAG rabbit polyclonal (sc-807, Santa Cruz); anti-HA mouse monoclonal (clone 12CA5, Boehringer Mannheim); anti-IL-1α and anti-TNFα rabbit polyclonal (BioTech Trade, St Leon-Rot); anti-IL-1 $\alpha$  rabbit monoclonal (clone #1190, a gift from Dainippon Pharmaceutical Co., Ltd, Japan); anti-IL-1a goat polyclonal (R&D systems); anti-IgG rabbit polyclonal (Dianova); anti-p53 mouse monoclonal (clone Ab-2, Calbiochem); anti-p50 antibody (sc-1190X, Santa Cruz); and the anti-p65 antibody (sc-372, Santa Cruz). Cells were incubated with 10 ng/ml human recombinant TNFa (Calbiochem), 10 ng/ml human recombinant IL-1α (Sigma) or 60 ng/ml TPA (Sigma). Pervanadate was prepared as previously described (Imbert et al., 1996) with freshly dissolved Na<sub>3</sub>VO<sub>4</sub> pH10 (Sigma). For UV irradiation, cells were washed with phosphate-buffered saline (PBS) and irradiated without PBS with a monochromatic UV lamp (15W Hg lamp) obtained from Vetter (Wiesloch, Germany) emitting a wavelength of 254 nm (halfmaximal width, 2.3 nm).

#### Nuclear extracts and electrophoretic mobility shift assay

Nuclear extracts were prepared and used in EMSAs as previously described (Stein *et al.*, 1989a). Briefly, cells were washed twice with ice-cold PBS, scraped from the plate with a rubber policeman and suspended in 100  $\mu$ l lysis buffer (10mM HEPES pH 7.9, 1 mM EDTA, 60 mM KCl, 0.5% Nonidet P-40, 1 mM DTT, 1 mM PMSF). After 5 min on ice, nuclei were sedimented at 1200 g for 5 min. The supernatant was diluted with 50  $\mu$ l of nuclear buffer (250 mM Tris–HCl pH 7.8, 60 mM KCl, 1 mM DTT, 1 mM PMSF), cleared by centrifugation at 13 000 g for 15 min and used as cytoplasmic extracts. The nuclei were washed with lysis buffer without NP-40 and suspended in 100  $\mu$ l nuclear buffer. Nuclei were lysed by three cycles of freezing and thawing in liquid nitrogen and ice. The nuclear extracts were cleared by centrifugation at 13 000 g for 15 min.

EMSAs were done as described previously (Stein *et al.*, 1989a), with slight modifications: binding was performed in a volume of 20  $\mu$ l with 5–6  $\mu$ g nuclear protein in a buffer containing 12 mM HEPES pH 7.8, 62.5 mM Tris–HCl pH 7.8, 60 mM KCl, 0.6 mM EDTA, 12% glycerol, 5 mM DTT, 2  $\mu$ g BSA and 1  $\mu$ g poly(dI–dC). Approximately 10 fmol (100 000 c.p.m.) of <sup>32</sup>P-radiolabeled double-stranded oligonucleotide (5'-AGCTTGGGGACTTTCCAGCCG-3') derived from the HIV-LTR, or (5'-TCGACTCGAGATGCAAATAAG-3') for Oct-1, were used per reaction. Resultant DNA–protein complexes were resolved on 5% polyacrylamide gels and detected by autoradiography.

#### Immunoprecipitation and Western blot analysis

Cells were lysed in a buffer containing 137 mM NaCl, 20 mM Tris pH 8.0, 2 mM EDTA, 1% Triton X-100, 10% glycerol, 50 mM NaF, 10 mM NaPPi, 10  $\mu$ M Na<sub>2</sub>MoO<sub>4</sub>, 50  $\mu$ M ZnCl<sub>2</sub>, 20 mM  $\beta$ -glycerolphosphate, 1 mM PMSF, 1 mM Na<sub>3</sub>VO<sub>4</sub>, 1  $\mu$ g/ml aprotinin and 10  $\mu$ g/ml leupeptin. To separate nuclear and cytoplasmic extracts the

Immunoprecipitations were performed by using 20  $\mu$ l protein A beads (Pharmacia) and 5  $\mu$ l of antibody in a volume of 800  $\mu$ l lysis buffer for 2 h at 4°C. The beads were pelleted and washed three times with lysis buffer and once with PBS before mixing with 45  $\mu$ l 2× SDS sample buffer. The samples were boiled for 5 min and resolved by SDS–polyacrylamide gel electrophoresis (PAGE), followed by Western blot analysis.

For Western blot analyses equal amounts of protein extracts were fractionated by SDS–PAGE and electrophoretically transferred to a polyvinylidene difluoride membrane (Millipore). Membranes were blocked (1 h at room temperature) with PBS containing 0.3% Tween 20 and 10% powdered milk and then incubated with a specific antiserum. Specific proteins were visualized by enhanced chemiluminescence (Amersham Life Science Inc., Cleveland, OH).

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