Differential interactions of Rel-NF- κ B complexes with $I\kappa$ B α determine pools of constitutive and inducible NF- κ B activity

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The Rel-NF-KB family of transcription factors plays a crucial role in the regulation of genes involved in inflammatory and immune responses. We demonstrate that in vivo, in contrast to the other members of the family, RelB associates efficiently only with NF- κ B1 (p105-p50) and NF-kB2 (p100-p52), but not with cRel or p65. The RelB-p52 heterodimers display a much lower affinity for $I \kappa B \alpha$ than RelB-p50 heterodimers or p65 complexes. However, similarly to the other Rel-NF-KB complexes, RelB-p52 can upregulate the synthesis of $I \kappa B \alpha$ leading to the cytoplasmic trapping of dimers which have a higher affinity for the inhibitor. We suggest that a hierarchy of interactions between IKBa and the different Rel-NF-KB complexes governs their cellular distribution. This results in the presence of two distinct pools of NF-KB activity which differ in their composition: one a constitutive nuclear and the other an inducible cytoplasmic activity.

Key words: gene regulation/I κ B α interaction/Rel-NF- κ B complex/transcription factor

Introduction

One of the central problems of molecular biology is to understand the mechanisms underlying temporal and tissue-specific gene expression and activation of specific genetic programs in response to extracellular stimuli. Accumulated evidence indicates that transcription initiation represents the major regulatory event controlling specificity of gene expression. This is carried out by the combined action of transcription factors interacting with the regulatory elements of genes. Rapid responses to external signals are often mediated by the inducible activation of transcription factors by post-translational mechanisms, as opposed to their de novo synthesis. Examples of post-translational mechanisms serving this purpose are the ligand-induced DNA binding of nuclear receptors (reviewed in Evans, 1988; Green and Chambon, 1988) and phosphorylation (reviewed in Hunter and Karin, 1992). Interactions with inhibitory molecules represent yet another mechanism to control the activity of transcription factors which would be amenable to rapid induction. Associations of NF- κ B proteins with the I κ B family of

inhibitors is the best studied example of such a controlling mechanism.

The Rel-NF- κ B family of transcription factors is involved in the regulation of immune and acute-phase responses at the transcriptional level (reviewed in Grilli et al., 1993; Liou and Baltimore, 1993). The mammalian members of this family include p105-p50 (NF-KB1, also known as KBF1 and H2TF1), p100-p52 (NF-κB2, also known as p50B, p49 and Lyt-10), cRel, p65 (RelA) and RelB. All members of the Rel-NF-kB family share a 300 amino acid region of homology called the Rel homology domain (RHD) which mediates dimerization and DNA binding (for a review see Baeuerle, 1991; Blank et al., 1992; Bose, 1992; Gilmore, 1992; Nolan and Baltimore, 1992; Grilli et al., 1993; Grimm and Baeuerle, 1993). The activity of Rel-NF-kB complexes is regulated by their interactions with the various IKB molecules (reviewed in Nolan and Baltimore, 1992; Beg and Baldwin, 1993; Gilmore and Morin, 1993). Association with IkBa results in the cytoplasmic retention of NF-KB dimers which form an inactive, inducible cytoplasmic pool of NF- κ B. NF- κ B activity, while constitutive in mature B cells, can be rapidly induced in other cells by a variety of agents including cytokines, pathogen infections or reactive oxygen intermediates (Grilli et al., 1993; Grimm and Baeuerle, 1993). The induction of Rel-NF-KB activity involves the release of $I\kappa B\alpha$ from the Rel-NF- κB complexes, allowing their rapid translocation to the nucleus (reviewed in Beg and Baldwin, 1993; Gilmore and Morin, 1993). The biochemical mechanism underlying this dissociation has not been fully elucidated yet, but available evidence indicates that it involves $I\kappa B\alpha$ phosphorylation (Cordle et al., 1993; Devary et al., 1993; Diaz-Meco et al., 1993; Finco and Baldwin, 1993; Li and Sedivy, 1993) and degradation (Beg et al., 1993; Brown et al., 1993; Cordle et al., 1993; Henkel et al., 1993; Rice and Ernst, 1993; Sun et al., 1993; Chiao et al., 1994; Frantz et al., 1994). The precursors of p50 and p52, p105 and p100 respectively, show significant homology to the IkB molecules in their C-terminal region, and it has been suggested that they also trap NF- κB complexes in the cytoplasm (Inoue et al., 1992; Rice et al., 1992; Mercurio et al., 1993; Naumann et al., 1993; Scheinman et al., 1993; Sun et al., 1994).

It has been demonstrated recently that p65- and cRelcontaining complexes induce expression of $I\kappa B\alpha$, creating an autoregulatory loop in which increased synthesis of inhibitor serves to restore the uninduced state of the cell (Brown *et al.*, 1993; de Martin *et al.*, 1993; Scott *et al.*, 1993; Sun *et al.*, 1993). Such an autoregulatory mechanism would ensure the transient nature of the NF- κ B-mediated gene activation. Thus, one way to regulate the NF- κ Bmediated genetic response is by controlling the cellular localization of the dimers. Rapidly accumulating evidence suggests that the various Rel–NF- κ B complexes have different target specificities and that levels of transcriptional activation elicited by these transactivators differ depending on the particular κ B binding site, despite comparable levels of DNA binding (Schmid *et al.*, 1991; Fujita *et al.*, 1992; Hansen *et al.*, 1992; Kretzschmar *et al.*, 1992; Kunsch *et al.*, 1992; Nakayama *et al.*, 1992; Perkins *et al.*, 1992; Tan *et al.*, 1992; Sica *et al.*, 1992; Kunsch and Rosen, 1993; Narayanan *et al.*, 1993; Shu *et al.*, 1993; Hansen *et al.*, 1994). These data thus underscore the importance of individual complexes and indicate that combinatorial associations of Rel–NF- κ B proteins provide another way of controlling the specificity of the NF- κ B-mediated responses.

We have been interested in understanding the physiological role of RelB (Ryseck et al., 1992; Carrasco et al., 1993; Dobrzanski et al., 1993). In this report we analyse the capacity of RelB to form complexes with other members of the family and the interactions of RelB heterodimers with $I\kappa B\alpha$ —the prototype of an $I\kappa B$ inhibitor. We have found that in vivo, RelB, in contrast to p65 and cRel, forms dimers only with p50, p52 and their respective precursors p105 and p100. Interestingly, the RelB-p52 heterodimer does not associate efficiently with I κ B α ; however, similarly to the other Rel-NF- κ B complexes, RelB heterodimers can upregulate the levels of endogenous I κ B α . Our data indicate that this leads to the cytoplasmic trapping of Rel-NF-kB complexes with a higher affinity for the inhibitor, like the p65-containing complexes, thus changing the composition of the NF-KB activity in the cell. We demonstrate further that $I\kappa B\alpha$ has a differential affinity for the various Rel-NF-KB complexes in vivo, suggesting a hierarchy of interactions which determines the cellular distribution of Rel-NF-KB dimers. Based on these results we propose a model suggesting how the specificity of action of the NF-KB activity can be achieved given the large number of possible combinations present in the cell.

Results

RelB is involved in a limited number of interactions in vivo

To determine the capacity of RelB to form complexes with other Rel-NF- κ B proteins in vivo, T and B cell lines were screened for the expression of RelB. Daudi cells (a human B cell line) were chosen for further analysis because they express high levels of RelB and other family members (Figure 1A). To identify which members of the Rel-NF-KB family interact with RelB, Daudi cells were labeled with [35S]methionine, and proteins associated with RelB were screened using specific antibodies. The results indicate that RelB forms heterodimers with p52 and p50, and with their respective precursors p100 and p105 (Figure 1B). The amount of p100-p52 and p105-p50 associated with RelB is similar to that found with p65 and cRel (Figure 1C and D). Since RelB, p65 and cRel are expressed at comparable levels, the results suggest that their capacities to interact with p100-p52 and p105-p50 are similar. However, in contrast to p65 which readily forms complexes with cRel (Figure 1C and D; Hansen et al., 1992, 1994), RelB interacts with cRel only weakly, if at all (Figure 1B and D). No significant interaction between



Fig. 1. Analysis of Rel–NF-κB complexes in Daudi cells. Daudi cells were labelled with [³⁵S]methionine and lysed under denaturing (**A**) or native (**B**–**D**) conditions. To compare the ability of RelB, p65 and cRel to form complexes *in vivo*, native cell lysates were incubated with the corresponding antibody. Immunocomplexes were precipitated with protein A–Sepharose. The immunoprecipitated proteins were released, sequentially reprecipitated with the indicated antibody and subjected to SDS–PAGE, followed by fluorography. All immunoprecipitations were performed as described in Materials and methods. (A) Total amounts of Rel–NF-κB proteins present in Daudi cells. Each protein was immunoprecipitated twice with the specific antibody. (B) Analysis of RelB-containing complexes. (C) Analysis of p65-containing complexes. (D) Analysis of cRel-containing complexes.

RelB and p65 was detected *in vivo* (Figure 1B and C), in accordance with our *in vitro* results (R.-P.Ryseck and P.Dobrzanski, unpublished results). These data demonstrate that RelB has a more restricted ability to form dimers compared with the other members of the family.

RelB–p52 heterodimer is not inhibited by any of the known IkB molecules

Since the activity of the members of the Rel–NF- κ B family of transcription factors is regulated by their interaction with the I κ B family of molecules (for a review see Nolan and Baltimore, 1992; Beg and Baldwin, 1993; Gilmore and Morin, 1993), we analyzed whether RelBcontaining heterodimers are associated with I κ B α *in vivo*. Surprisingly, we found very little if any I κ B α co-immunoprecipitated with RelB complexes (Figure 1B), in contrast to those containing p65 or cRel which are associated with significant amounts of I κ B α (Figure 1C and D). This result suggests that RelB–p52, which as shown above represents the major RelB-containing heterodimer in Daudi cells, does not interact efficiently with I κ B α . Our previous *in vitro* observation that the DNA binding of RelB–p50 complexes can be blocked by I κ B α (Tewari *et al.*, 1992),

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Fig. 2. Effect of different $I\kappa B$ molecules on the transcriptional activity of RelB-p52 and RelB-p50 heterodimers. Jurkat cells were cotransfected with expression vectors encoding RelB and p52, or RelB and p50, together with vectors expressing $I\kappa B\alpha$, $I\kappa B\gamma$ or Bcl3, as indicated, and the $8 \times \kappa B$ -tk-CAT reporter plasmid. Cells were harvested 48 h after transfection. Transfections and CAT assays were performed as described in Materials and methods. (A) Effect of different $I\kappa B$ proteins on RelB-p52 and RelB-p50 activity. Equal amounts (3 µg) of each of the expression vectors were cotransfected with 2 µg of the reporter plasmid. The total amount of DNA transfected was adjusted to 18 µg with the expression vector pMexneo. Upper panel: transcriptional activity was calculated as an average of three independent experiments quantitated on a PhosphorImager. The expression generated by the cotransfection experiment. Transfection of pMexneo alone serves as a control (C). (B) RelB-p52 and RelB-p50 are differentially inhibited by $I\kappa B\alpha$. Constant amounts (2.5 µg) of each of RelB-, p52- and p50-expressing vectors were cotransfected with increasing amounts, as indicated, of vector expressing $I\kappa B\alpha$ and 2 µg of the reporter plasmid. Details of upper and lower panels are as described in (A).

prompted us to test whether RelB-p52 DNA binding can also be inhibited by I κ B α . In agreement with the weak interactions observed *in vivo*, *in vitro*-translated RelB-p52 heterodimers consistently required 2- to 4-fold more I κ B α than RelB-p50 heterodimers for the full inhibition of DNA binding (not shown).

To determine whether the observed differential inhibition by IkBa of the DNA binding of RelB complexes in vitro is reflected in a differential repression of their transcriptional activity, transient transfection assays were performed. Jurkat cells were cotransfected with equimolar amounts of expression vectors coding for RelB, IkBa and either p52 or p50, together with a reporter construct carrying the chloramphenicol acetyltransferase (CAT) gene under the control of κB binding sites. The cotransfection of equimolar amounts of vector encoding IkBa results in a strong inhibition of the RelB-p50 transcriptional activity (down to 11% of the control without inhibitor), whereas the activity of RelB-p52 is only slightly reduced (Figure 2A). Therefore, it was of interest to determine whether the transcriptional activity of RelB-p52 heterodimers could be efficiently inhibited by other members of the

IkB family, namely IkB γ (Hatada *et al.*, 1992; Inoue *et al.*, 1992; Liou *et al.*, 1992) and Bcl3 (Ohno *et al.*, 1990; Kerr *et al.*, 1991; Nolan *et al.*, 1993). To this end, RelB and p52 or RelB and p50, were cotransfected together with equimolar amounts of vectors coding for IkB γ or Bcl3. As demonstrated in Figure 2A, IkB γ showed no effect and Bcl3 only minimal inhibition of the transcriptional activity of both RelB heterodimers. Western blot analysis verified that all vectors expressed comparable amounts of protein (not shown).

To analyse further the differential effect of $I\kappa B\alpha$ on the transcriptional activity of RelB-p52 and RelB-p50 heterodimers, constant amounts of activators were cotransfected with increasing amounts of $I\kappa B\alpha$. As shown in Figure 2B, four times more $I\kappa B\alpha$ was required to significantly inhibit RelB-p52 activity (56%) compared with RelB-p50 (60%). The activity of RelB-p50 was strongly inhibited by equimolar amounts of $I\kappa B\alpha$ expression vector, whereas a 2-fold molar excess basically eliminated all transcriptional activity. In contrast, a 4-fold excess of the $I\kappa B\alpha$ expression vector was required to completely inhibit the transcriptional activity of



Fig. 3. Differential effect of $I\kappa B\alpha$ on RelB-p52 and RelB-p50 DNA binding activity *in vivo*. COS cells were transfected with constant amounts (2 µg each) of vectors expressing RelB and p52 or RelB and p50, together with increasing amounts of $I\kappa B\alpha$ expressing vector (as indicated). The total amount of DNA transfected was adjusted to 10 µg with expression vector. Whole-cell extracts were prepared 48 h after transfection. (A) EMSA with extracts prepared from cells transfected with the indicated expression vectors. To confirm the identity of the extract-generated bands, specific antibodies were included in binding reactions, as shown. Extract from cells transfected with the expression vector alone served as a control (C). (B) Quantitation of EMSA results. An average of two independent experiments quantitated with a PhosphorImager is shown. The values obtained for heterodimers were normalized to the amount of p52 or p50 homodimers binding, and serve as an internal control. Binding of the heterodimer without cotransfected IkB\alpha represents 100% activity. (C) Expression of RelB, p52, p50 and IkB\alpha in transfected COS cells. Parallel whole-cell extracts used for EMSA (A) were subjected to SDS-PAGE, followed by Western blotting analysis using specific antibodies, as indicated. Proteins were detected using the ECL system.

RelB-p52, further confirming that $I\kappa B\alpha$ has a lower affinity for RelB-p52 than for RelB-p50. These results indicate that *in vivo* RelB-p52 is not efficiently inhibited by any of the known inhibitors (I $\kappa B\alpha$, I $\kappa B\gamma$ and Bcl3), whereas the activity of RelB-p50 can be modulated only by I $\kappa B\alpha$.

To investigate whether the observed differential inhibition of transcriptional activity of RelB-p52 versus RelB-p50 could be correlated with the inhibition of DNA binding, COS cells were transfected with vectors that express high levels of RelB, p52, p50 and IkBa. Constant amounts of RelB and p52 or RelB and p50, were cotransfected together with increasing amounts of IkBa. Wholecell extracts were prepared from transfected cells and kB binding activity was determined in electrophoretic mobility shift assays (EMSA; Figure 3A). The control lane showed very little binding activity, whereas extracts from transfected cells generated two major bands. As determined by specific antibodies, the upper band corresponds to either the RelB-p52 or RelB-p50 heterodimer, and the lower band represents either the p52 or p50 homodimer. The anti-p50 antibody completely abolished binding of the p50 homodimer and reduced, but did not eliminate, binding of the RelB-p50 heterodimer, in accordance with previous reports demonstrating that the p50-specific antiserum reacts better with homodimers (Kieran et al., 1990; Lernbecher et al., 1993). Even at the lowest amount of transfected I κ B α , the binding of the RelB-p50 complex is strongly reduced. The highest amount of inhibitor transfected almost abolished binding of RelB-p50, whereas RelB-p52 still displayed significant binding. The binding activity of the homodimers was, as expected, not affected by the increasing amounts of $I\kappa B\alpha$ expressed and served as an internal control. Figure 3B shows the quantitation of two independent experiments. The data demonstrate that *in vivo* the DNA binding activity of RelB-p52 is much more resistant to inhibition by $I\kappa B\alpha$ than is the binding of RelB-p50. Western blot analysis verified that comparable amounts of protein were expressed in transfected cells (Figure 3C). These data validate our interpretation of the EMSA results which are consistent with those obtained in the CAT assays (Figure 2B).

RelB heterodimers increase levels of endogenous $l\kappa B \alpha$

The lack of efficient inhibition of RelB-p52 activity by any of the known IkB molecules in vivo was particularly intriguing in view of the recent observations that other members of the Rel-NF- κB family, namely p65 and cRel, induce expression of $I\kappa B\alpha$, generating an autoregulatory feedback mechanism (Brown et al., 1993; de Martin et al., 1993; Scott et al., 1993; Sun et al., 1993). To test whether the RelB heterodimers can increase the level of endogenous IkBa, COS cells were transfected with vectors expressing RelB and p52, RelB and p50 or p65 alone. After 48 h the level of endogenous I κ B α was analyzed by Western blot (Figure 4A). RelB-p52 and RelB-p50 strongly increase the cellular level of $I\kappa B\alpha$. We consistently observed higher levels of IkBa protein following transfection of p65 than of RelB heterodimers. This is probably due to the increased stability of IkBa resulting from its stronger interaction with p65 than with RelB-p52 or RelB-p50. Nevertheless, the results indicate that RelB-p52 and RelB-p50 strongly induce expression of IkBa.



Fig. 4. (A) RelB heterodimers increase the levels of endogenous $I\kappaB\alpha$. COS cells were transfected with expression vectors encoding the indicated Rel-NF- κ B proteins. As a control, the expression vector alone was transfected (C). Whole-cell extracts prepared 48 h after transfection were subjected to SDS-PAGE, followed by Western blot analysis using the ECL system. (B) Immunofluorescence analysis of the cellular localization of p65 and RelB following cotransfection of vectors expressing p65, RelB and p52 into COS cells. Fixed cells were subjected to immunofluorescence using anti-RelB- (left panel) or anti-p65 (right panel)-specific antibodies. FITC-conjugated donkey antibody against rabbit IgG was used as a secondary antibody. Similar results were obtained for p65, RelB and p50 cotransfections (not shown). (C) IkB α differentially interacts with p65 and RelB complexes *in vivo*. COS cells were transfected with p65, RelB and p52 (left panel) or p65, RelB and p50 (right panel). Forty-eight h later, cells were labeled with [³⁵S]methionine and lysed under native conditions. Extracts were incubated sequentially with antibodies against p65, RelB and IkB α . The immunoprecipitated complexes were released and tested for the presence of IkB α using specific antibodies. Lanes p65 and RelB indicate the IkB α found associated with p65 and RelB, respectively. IkB α not associated with p65 or RelB, i.e. what remains in the extract following depletion of p65 and RelB complexes by the corresponding antibodies, is referred to as 'free'. All incubations with antibodies and protein A-Sepharose were performed for 2 h at 4°C. (D) Effects of IkB α on the transcriptional activity of RelB-p52, RelB-p50 and p65-p50. Jurkat cells were cotransfected with a constant amount (3 μ g) of each of RelB-, p65-, p52- and p50-expressing vectors, together with the indicated amount of IkB α expressing vector and 2 μ g of the reporter plasmid. The total amount of transfected DNA was adjusted to 12 μ g with expression vector. All other d

RelB heterodimers represent a constitutive NF- κ B activity

The induction of I κ B α by RelB-p52, together with its low affinity for the inhibitor, suggested that RelB-p52 might be part of a constitutive nuclear NF-kB activity. Its ability to upregulate $I\kappa B\alpha$ synthesis would ensure cytoplasmic trapping of p65-containing complexes. This assumption was tested in triple cotransfection experiments in which COS cells were cotransfected with vectors expressing p65, RelB and p52 or p65, RelB and p50. After 48 h the cellular localization of the transfected proteins was determined by immunofluorescence and Western blot analyses of the nuclear and cytoplasmic extracts. Immunofluorescence assays demonstrate (Figure 4B) that in $\sim 90\%$ of expressing cells p65 was exclusively cytoplasmic, whereas 10% of cells showed staining in both cytoplasm and nucleus. In contrast, RelB staining was found mainly in the nucleus. By Western blot analysis we further verified that RelB, p52 and p50 were predominantly nuclear, whereas the majority of p65 was found in the cytoplasmic fraction (not shown). In control experiments RelB-p52 and RelB-p50 alone were transfected into COS cells and their cellular localization was analysed by immunofluorescence and Western blotting of nuclear and cytoplasmic fractions. We found that both heterodimers were predominantly nuclear, suggesting that weak interactions of RelB complexes with IKBa are not limited to lymphoid cells (not shown). To ensure that the majority of the RelB molecules are complexed, cotransfections were performed with a relative excess of p52 or p50, so that the observed nuclear localization of RelB is not due to free, uncomplexed RelB migrating to the nucleus, but reflects the true cellular distribution of RelB-p52 and RelB-p50. The difference in the cellular localization of p65- and RelB-containing complexes suggests that RelB-p52 represents the constitutive NFκB activity, in contrast to p65 complexes which remain in the cytoplasm, probably due to their association with the endogenous I κ B α . Interestingly, under these conditions RelB-p50 behaves like RelB-p52 and is found predominantly in the nucleus (not shown). This indicates that the affinity of RelB-p50 for I κ B α , although higher than that of RelB-p52, is still lower than that of p65 complexes, suggesting that in vivo in the presence of limiting amounts of $I\kappa B\alpha$, RelB-p50 might in part contribute to the constitutive NF-kB activity.

To test whether the cytoplasmic localization of p65 in transfected COS cells was due to its association with endogenous $I\kappa B\alpha$, the amount of $I\kappa B\alpha$ associated with p65 and RelB was estimated by immunoprecipitation following [³⁵S]methionine labeling of the cells. Figure 4C demonstrates that ~90% of the total cellular $I\kappa B\alpha$ is found associated with p65, whereas only a minor fraction interacts with RelB-p52 or RelB-p50 heterodimers. The absence of free $I\kappa B\alpha$ indicates that the inhibitor is present in limiting quantities. Since p65 and RelB were expressed at similar levels (not shown), the observed differences in the amount of $I\kappa B\alpha$ associated with these complexes



Fig. 5. Analysis of Rel–NF-κB binding activity in Daudi cells. Whole-cell extracts were prepared from Daudi cells which were either unstimulated or incubated for 45 min with 100 ng/ml of PMA. κB binding activity was tested in EMSA using a ³²P-labeled palindromic κB probe. To analyze the identity of DNA binding complexes, specific antibodies were included in the binding assays as indicated. Equal amounts of a preimmune serum were included in control reactions (PI).

reflects the real differences in their affinity for $I\kappa B\alpha$. These results suggest a hierarchy of interactions between IκBα and different RelB and p65 complexes. To compare more directly the affinity of $I\kappa B\alpha$ for RelB-p52, RelB-p50 and p65-p50, transient transfection assays were performed. Jurkat cells were cotransfected with constant amounts of RelB-p52, RelB-p50 and p65-p50 together with increasing amounts of inhibitor. Figure 4D demonstrates that transcriptional activity of p65-p50 heterodimer is indeed the most sensitive to the IkBamediated inhibition, followed by that of RelB-p50. These data further confirm that $I\kappa B\alpha$ can differentially interact with various Rel-NF-κB complexes. Thus, in vivo, IκBα has much higher affinity for p65 complexes than for any of the RelB heterodimers, providing an explanation for the cytoplasmic localization of p65 in cotransfection assays.

Our results further suggest that in unstimulated cells expressing RelB, p50, p52 and p65, RelB complexes will constitute the major nuclear κB binding activity, whereas the p65 complexes would be associated with $I\kappa B\alpha$ and therefore be retained in the cytoplasm. To verify these predictions, the NF-kB binding activity present in Daudi cells was analyzed by EMSA. The extract from unstimulated cells generated a major band which contained RelB but not p65, as demonstrated by its sensitivity to the corresponding antibodies (Figure 5), indicating that in unstimulated cells there is very limited, if any, κB binding activity which could be attributed to p65 complexes. Activation of cells with phorbol 12-myristate 13-acetate (PMA) induces a new binding activity which contains p65, as demonstrated by its abolition by the corresponding antibody. In contrast, DNA binding of RelB heterodimers is not increased by treatment with PMA, suggesting that they are constitutively active in the cell. PMA treatment has been shown to induce κB binding activity by releasing Rel-NF- κ B dimers from their association with I κ B α (Henkel et al., 1993; Sun et al., 1993). The data thus demonstrate that in cells expressing RelB and p65 the latter remains complexed with $I\kappa B\alpha$ and as a consequence represents the inducible NF- κ B activity, while RelB complexes represent the constitutive activity.

Discussion

Interactions of ReIB with other members of the ReI–NF- κ B family

We have determined the capacity of RelB to interact with the other members of the NF-kB family in vivo. Our studies demonstrate that RelB forms heterodimers only with p50 and p52 and their respective precursors p105 and p100. The amounts of p50-p105 and p52-p100associated with RelB reflect the relative abundance of these proteins in the cell, suggesting that RelB has similar affinity for these molecules. We also found that p65 and cRel associate with p52-p100 and p50-p105 with similar affinities to RelB. These results led us to conclude that the interactions of RelB and p65 with p52-100 and p50-105 are governed by the concentrations of the relevant proteins in the cell. However, in contrast to p65 which readily forms complexes with cRel (Hansen et al., 1992, 1994), we found no evidence of RelB-cRel interactions in vivo, in accordance with our in vitro observations. We also failed to detect any significant interactions between RelB and p65. These data, together with the inability of RelB to form homodimers (Ryseck et al., 1994), indicate that RelB, compared with the other members of the family, is involved in a limited number of interactions in vivo. Such a restricted ability to form dimers might suggest that RelB plays a specific function in regulating the expression of a particular set of genes in vivo.

Since RelB-p50 and RelB-p52 heterodimers are strong transactivators (Bours et al., 1992; Ryseck et al., 1992), we analyzed if this activity can be modulated by members of the I κ B family. We found that I κ B α displayed a differential inhibition on both complexes. Several lines of evidence, including EMSA of proteins generated in vitro and in vivo, and transient transfection assays demonstrated that RelB-p52 has much lower affinity for I κ B α than RelB-p50. However, transcriptional activity of RelBp52 can be fully inhibited by an excess of IkBa (Figure 2). In view of the recent findings that the uncomplexed IkB α is degraded rapidly in the cells (Beg et al., 1993; Brown et al., 1993; Henkel et al., 1993; Rice and Ernst, 1993; Sun et al., 1993; Chiao et al., 1994; Frantz et al., 1994), it seems unlikely that such an excess of $I\kappa B\alpha$ can be achieved in vivo. The lack of interaction between RelB-p52 and I κ B α in Daudi cells, despite the abundance of the inhibitor, supports the notion that $I\kappa B\alpha$ is a weak modulator, if at all, of RelB-p52 activity in vivo. Interestingly, we have also found that none of the other known IkB molecules, IkBy or Bcl3 substantially inhibit the activity of RelB-p52 or RelB-p50. The lack of efficient in vivo inhibition of the RelB-p52 heterodimer by any of the known $I\kappa B$ molecules suggests that this complex might function as a constitutively active transcription factor. These data emphasize the importance of the processing of the precursors p100 and p105 as a regulatory event controlling NF-KB activity (Inoue et al., 1992; Rice et al., 1992; Mercurio et al., 1993; Naumann et al., 1993; Scheinman et al., 1993; Sun et al., 1994).

One might expect, however, that such a highly regulated system as NF- κ B – I κ B should have the ability to modulate



Fig. 6. Model explaining how the differential interactions with $I\kappa B\alpha$ determine the constitutive versus inducible character of Rel-NF-κB activity leading to the existence of two pools of complexes which differ in their composition and cellular localization. 'X' represents a subset of NF-κB-regulated genes driven by the constitutively active RelB-p52 dimers. In response to the external signal the inducible pool, represented here by p65 complexes, triggers a specific genetic program (Y). The dashed circle is any member of the Rel-NF-κB family which combines with p65 to form dimers and with high affinity for IκBα. Cytoplasm (C) and nucleus (N) are indicated. See text for a discussion.

the activity of RelB-p52. It is then plausible that in certain cells RelB-p52 activity could be regulated by an as yet unknown inhibitor. One might also consider other possible regulatory mechanisms such as post-transcriptional modification or modulatory interactions with other proteins.

Differential interactions of Rel–NF- κ B complexes with $l\kappa$ B α and their functional consequences

The lack of efficient inhibition of the RelB-p52 activity by any of the known IKB molecules in vivo was of particular interest considering the recent reports which show that other members of the Rel-NF- κ B family, p65 and cRel, induce expression of IkBa, generating an autoregulatory feedback mechanism (Brown et al., 1993; de Martin et al., 1993; Scott et al., 1993; Sun et al., 1993). The findings that various Rel-NF-KB complexes, including both RelB heterodimers, are able to upregulate the synthesis of $I\kappa B\alpha$, together with our observation that these complexes exhibit differential affinity for IkBa (Figure 4D), have an important implication for the regulation of NF-KB activity. It suggests a model (Figure 6) which predicts that RelB-p52 would function as a constitutive nuclear NF-kB activity, regulating expression of specific genes, including IkBa. This would ensure a constant supply of $I\kappa B\alpha$ to the cytoplasm, trapping NF- κB molecules with high affinity for I $\kappa B\alpha$, like p65-p50 or p65-p52. These complexes would then represent the inactive cytoplasmic reservoir of the inducible NF-KB activity. The results of the triple transfection experiments (Figure 4B) and the analysis of the constitutive and PMAinducible NF-kB activity in extracts from Daudi cells (Figure 5) fully support this model. In addition, RelB, cRel and p65 complexed with precursors p100 and p105 are also part of the cytoplasmic fraction. However, since

the proteolytic processing, even after activation, is rather slow (Mercurio *et al.*, 1993), these complexes would not contribute to the rapid alteration of the NF- κ B activity in response to the changing environment.

Our model predicts that the constitutive nuclear and the inducible cytoplasmic pools would consist of different Rel-NF-KB complexes, e.g. RelB-p52 versus p65-containing complexes. This is important in view of the rapidly accumulating evidence indicating that different Rel-NFκB complexes exhibit vastly different target specificities (Schmid et al., 1991; Fujita et al., 1992; Hansen et al., 1992; Kretzschmar et al., 1992; Kunsch et al., 1992; Nakayama et al., 1992; Perkins et al., 1992; Sica et al., 1992; Tan et al., 1992; Kunsch and Rosen, 1993; Narayanan et al., 1993; Shu et al., 1993). Therefore, the differential regulation of the various complexes would enable the cell to turn on specific genetic programs in response to external stimuli by activating a specific set of Rel-NF- κ B complexes. One could further predict that the inducible activity would activate genes involved in rapid responses like the acute-phase response and inflammation. The constitutive activity would regulate those genes whose protein products are required for a prolonged period of time, like immunoglobulin G or major histocompatibility complex. According to the model, the predicted cellular distribution of the complexes would be determined by their relative affinity for $I\kappa B\alpha$ and by the relative abundance of the inhibitor and of NF-κB complexes. Those with the highest affinity for $I\kappa B\alpha$, like p65 heterodimers, would be mainly cytoplasmic and would represent the inducible pool. On the other hand, complexes with the lowest affinity for $I\kappa B\alpha$, like RelB-p52, would be predominantly nuclear and provide the constitutive activity. Furthermore, one could envisage that the constitutive versus inducible nature of certain complexes, like RelB-p50, would depend on the context of the Rel-NF- κB complexes expressed in the cell. In the presence of only RelB-p52 complexes, RelB-p50 would be a primary target for the cytoplasmic trapping by $I\kappa B\alpha$. However, in the presence of p65-p50 heterodimers the latter would mainly associate with the inhibitor, allowing RelB-p50 complexes to move to the nucleus. This would provide the system with the needed flexibility and would assure its dynamic nature. In a recent report, Lernbecher et al. (1993), using transgenic mice, correlated the expression of a reporter construct driven by κB binding sites with the presence of constitutively active RelB complexes in lymphoid tissues, supporting our model in an animal system. Although initially this concept was developed based on the experiments involving RelB and p65 heterodimers only, it can easily incorporate the cRel-containing complexes. It remains to be tested how the affinity of cRel complexes for IkBa compares in vivo with that of other Rel-NF-KB complexes. However, the recent observation by Liou and Baltimore (1993), who found that in certain B cell lines cRel-p50 is constitutively active, whereas p65-p50 is inducible, further supports our model and indicates that p65 heterodimers might indeed represent the 'ultimate' NF-kB-inducible activity. The very tight regulation of the p65-p50 activity suggests the importance of this heterodimer in cellular responses. Our model does not take into account the possible contribution of homodimers like p50-p50 or complexes between

p50 and Bcl3, since there is some controversy as to what the role of these complexes might be (Franzoso *et al.*, 1992; Fujita *et al.*, 1992). The existence of the nuclear and cytoplasmic pools of NF- κ B complexes, which differ in composition, explains at least in part the specificity of action that can be achieved given the large number of possible NF- κ B complexes.

Materials and methods

Cell culture and transfections

Jurkat cells were grown, electroporated and processed as described previously (Dobrzanski *et al.*, 1993). An $8 \times \kappa B$ -tk-CAT reporter vector was used in all experiments. CAT assays were performed as described (Gorman *et al.*, 1982). Reactions were normalized for β -galactosidase and/or protein concentrations. All CAT assays were quantitated in a PhosphorImager (Molecular Dynamics).

COS cells were transfected by DEAE-dextran methods as described (Seed and Aruffo, 1987). Typically, a 10 cm dish was transfected with $1-5 \mu g$ of each expression vector, as indicated in the figure legends. The total amount of DNA was adjusted with expression vector alone.

Antibodies

Specific antibodies used in these studies have been described previously (Ryseck *et al.*, 1992; Tewari *et al.*, 1992). The regions used for generating the fusion proteins were: for the murine cRel, amino acids 1–185 and 366–588; for RelB, amino acids 348–538; for p65, amino acids 399–489; for the N-terminal region of NF- κ B1, amino acids 1–120 (p50 subunit); for the C-terminal region of NF- κ B1, amino acids 575–972 (p105); for the N-terminal region of NF- κ B2, amino acids 775–901 (p100); and for the rat I κ B α , the complete coding sequence.

Labeling and immunoprecipitations

COS cells were rinsed twice with DMEM lacking methionine 48 h after transfection and then labeled for 2 h with [35 S]methionine (0.4 mCi/ml) in methionine-free medium.

For the labeling of Daudi cells, 2×10^6 cells were rinsed twice with DMEM lacking methionine and then labeled for 24 h with 1 mCi of [³⁵S]methionine/ml in DMEM containing 1 mg/l methionine supplemented with 20% dialyzed fetal calf serum. Immunoprecipitations under denaturing and native conditions were performed as described previously (Kovary and Bravo, 1991).

Immunofluorescence

COS cells were transfected with the indicated expression vectors. A total of 14–16 h after transfection, cells were trypsinized and plated onto coverslips. After a further 24–36 h, cells were fixed with methanol (2 min, room temperature) and incubated with the indicated antibody. FITC-conjugated donkey antibody against rabbit IgG was used as a secondary antibody.

Cell extracts and electrophoretic mobility shift assay

Whole-cell extracts from transfected COS cells were prepared 42–48 h after transfections, as described (Ganchi *et al.*, 1992). Whole-cell extracts from Daudi cells were prepared as described (Lernbecher *et al.*, 1993). κ B binding activity was analyzed in EMSA using a ³²P-labeled oligonucleotide containing a palindromic κ B site (Ballard *et al.*, 1990) as described (Tewari *et al.*, 1992), except that NP-40 was omitted in the binding reactions. Where indicated, specific antibodies were included.

Constructs

The CAT reporter construct $8 \times \kappa B$ -tk-CAT containing eight copies of the murine immunoglobulin enhancer κB site was generated as described (Ryseck *et al.*, 1992). The pMexneo (Martin-Zanca *et al.*, 1989)-based expression vectors for RelB and p50 were described (Ryseck *et al.*, 1992). Constructs expressing rat IkB α (Tewari *et al.*, 1992), human p52 (Bours *et al.*, 1992), IkB γ (Inoue *et al.*, 1992) and Bcl3 (Ohno *et al.*, 1990) were generated by inserting the corresponding coding sequences into pMexneo.

For experiments in COS cells the corresponding coding sequences were recloned into the expression vector pcDNAI/Amp (Invitrogen), except for the p50-expressing construct, Rc/CMV FLAGp50, which was a kind gift from Dr M.Karin.

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References

- Baeuerle, P.A. (1991) Biochim. Biophys. Acta, 1072, 63-80.
- Ballard,D.W., Walker,W.H., Doerre,S., Sista,P., Molitor,J.A., Dixon,E.P., Peffer,N.J., Hannink,M. and Greene,W.C. (1990) *Cell*, **63**, 803–814.
- Beg,A.A. and Baldwin,A.S., Jr (1993) Genes Dev., 7, 2064-2070.
- Beg,A.A., Finco,T.S., Nantermet,P.V. and Baldwin,A.S.,Jr (1993) *Mol. Cell. Biol.*, **13**, 3301–3310.
- Blank, V., Kourilsky, P. and Israel, A. (1992) Trends Biochem. Sci., 17, 135-140.
- Bose, H.R. (1992) Biochim. Biophys. Acta, 1114, 1-17.
- Bours, V., Burd, P.R., Brown, K., Villalobos, J., Park, S., Ryseck, R.-P., Bravo, R., Kelly, K. and Siebenlist, U. (1992) *Mol. Cell. Biol.*, **12**, 343–350.
- Brown, K., Park, S., Kanno, T., Franzoso, G. and Siebenlist, U. (1993) Proc. Natl Acad. Sci. USA, 90, 2532–2536.
- Carrasco, D., Ryseck, R.-P. and Bravo, R. (1993) Development, 118, 1221-1231.
- Chiao, P.J., Miyamoto, S. and Verma, I.M. (1994) Proc. Natl Acad. Sci. USA, 91, 28-32.
- Cordle,S.R., Donald,R., Read,M.A. and Hawiger,J. (1993) J. Biol. Chem., 268, 11803–11810.
- de Martin, R., Vanhove, B., Cheng, Q., Hofer, E., Csizmadia, V., Winkler, H. and Bach, F.H. (1993) *EMBO J.*, **12**, 2773–2779.
- Devary, Y., Rosette, C., DiDonato, J.A. and Karin, M. (1993) Science, 261, 1442–1445.
- Diaz-Meco, M.T. et al. (1993) Mol. Cell. Biol., 13, 4770-4775.
- Dobrzanski, P., Ryseck, R.-P. and Bravo, R. (1993) Mol. Cell. Biol., 13, 1572-1582.
- Evans, R.M. (1988) Science, 52, 889-895.
- Finco, T.S. and Baldwin, A.S., Jr (1993) J. Biol. Chem., 268, 17676-17679.
- Frantz,B., Nordby,E.C., Bren,G., Steffan,N., Paya,C.V., Kincaid,R.L., Tocci,M.J., O'Keefe,S.J. and O'Neill,E.A. (1994) *EMBO J.*, 13, 861–870.
- Franzoso, G., Bours, V., Park, S., Tomita-Yamaguchi, M., Kelly, K. and Siebenlist, U. (1992) Nature, 359, 339–342.
- Fujita, T., Nolan, G.P., Ghosh, S. and Baltimore, D. (1992) Genes Dev., 6, 775-787.
- Ganchi, P.A., Sun, S.-C., Greene, W.C. and Ballard, D.W. (1992) *Mol. Biol. Cell*, **3**, 1339–1352.
- Gilmore, T.D. (1992) Cancer Surv., 15, 69-87.
- Gilmore, T.D. and Morin, P.J. (1993) Trends Genet., 9, 427-433.
- Gorman, C., Moffat, L.F. and Howard, B. (1982) Mol. Cell. Biol., 2, 1044–1051.
- Green, S. and Chambon, P. (1988) Trends Genet., 4, 309-314.
- Grilli, M., Chiu, J.-S. and Lenardo, M.J. (1993) Int. Rev. Cytol., 143, 1-62.
- Grimm, S. and Baeuerle, P.A. (1993) Biochem. J., 290, 297-308.
- Hansen, S.K., Nerlov, C., Zabel, U., Verde, P., Johnsen, M., Baeuerle, P.A. and Blasi, F. (1992) *EMBO J.*, **11**, 205–213.
- Hansen,S.K., Baeuerle,P.A. and Blasi,F. (1994) Mol. Cell. Biol., 14, 2593–2603.
- Hatada,E.N., Nieters,A., Wulczyn,F.G., Naumann,M., Meyer,R., Nucifora,G., McKeithan,T.W. and Scheidereit,C. (1992) Proc. Natl Acad. Sci. USA, 89, 2489–2493.
- Henkel, T., Machleidt, T., Alkalay, I., Kronke, M., Ben-Neriah, Y. and Baeuerle, P.A. (1993) *Nature*, **365**, 182–185.
- Hunter, T. and Karin, M. (1992) Cell, 70, 375-387.
- Inoue, J.-I., Kerr, L.D., Kakizuka, A. and Verma, I.M. (1992) Cell, 68, 1109-1120.
- Kerr,L.D., Inoue,J., Davis,N., Link,E., Baeuerle,P.A., Bose,H.J. and Verma,I.M. (1991) Genes Dev., 5, 1464–1476.
- Kieran, M. et al. (1990) Cell, 62, 1007-1018.
- Kovary, K. and Bravo, R. (1991) Mol. Cell. Biol., 11, 2451-2459.
- Kretzschmar, M., Meisterernst, M., Scheidereit, C., Li, G. and Roeder, R.G. (1992) *Genes Dev.*, 6, 761–774.
- Kunsch, C. and Rosen, C.A. (1993) Mol. Cell. Biol., 13, 6137-6146.
- Kunsch,C., Ruben,S.M. and Rosen,C.A. (1992) Mol. Cell. Biol., 12, 4412-4421.
- Lernbecher, T., Muller, U. and Wirth, T. (1993) Nature, 365, 767-770.
- Li,S. and Sedivy,J.M. (1993) Proc. Natl Acad. Sci. USA, 90, 9247-9251.

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- Liou,H.-C. and Baltimore,D. (1993) Curr. Opin. Cell Biol., 5, 477-487.
- Liou,H.-C., Nolan,G.P., Ghosh,S., Fujita,T. and Baltimore,D. (1992) *EMBO J.*, **11**, 3003–3009.
- Martin-Zanca, D., Oskam, R., Mitra, G., Copeland, T. and Barbacid, M. (1989) Mol. Cell. Biol., 9, 24–33.
- Mercurio, F., DiDonato, J.A., Rosette, C. and Karin, M. (1993) *Genes Dev.*, **7**, 705–718.
- Nakayama,K., Shimizu,H., Mitomo,K., Watanabe,T., Okamoto,S.-I. and Yamamoto,K.-I. (1992) *Mol. Cell. Biol.*, **12**, 1736–1746.
- Narayanan, R., Higgins, K.A., Perez, J.R., Coleman, T.A. and Rosen, C.A. (1993) *Mol. Cell. Biol.*, **13**, 3802–3810.
- Naumann, M., Wulczyn, F.G. and Scheidereit, C. (1993) EMBO J., 12, 213-222.
- Nolan, G.P. and Baltimore, D. (1992) Curr. Opin. Genet. Dev., 2, 211-220.
- Nolan, G.P., Fujita, K., Bhatia, K., Huppi, C., Lio, H.-C., Scott, M.L. and Baltimore, D. (1993) *Mol. Cell. Biol.*, **13**, 3557–3566.
- Ohno, H., Takimoto, G. and McKeithan, T.W. (1990) Cell, 60, 991-997.
- Perkins, N.D., Schmid, R.M., Duckett, C.S., Leung, K., Rice, N.R. and Nabel, G.J. (1992) *Proc. Natl Acad. Sci. USA*, **89**, 1529–1533.
- Rice, N.R. and Ernst, M.K. (1993) EMBO J., 12, 4685-4695.
- Rice, N.R., MacKichan, M.L. and Israel, A. (1992) Cell, 71, 243-253.
- Ryseck, R.-P., Bull, P., Takamiya, M., Bours, V., Siebenlist, U., Dobrzanski, P. and Bravo, R. (1992) Mol. Cell. Biol., 12, 674–684.
- Ryseck, R.-P., Carrasco, D. and Bravo, R. (1994) In Tolle, T. and Zieglgansberger, W. (eds), *Immediate Early Genes: More Than Activity Marker*. Springer-Verlag, Berlin, Germany, in press.
- Scheinman, R.I., Beg, A.A. and Baldwin, A.S., Jr (1993) *Mol. Cell. Biol.*, **13**, 6089–6101.
- Schmid,R.M., Perkins,N.D., Duckett,C.S., Andrews,P.C. and Nabel,G.J. (1991) *Nature*, **352**, 733–736.
- Scott, M.L., Fujita, T., Lio, H.-C., Nolan, G.P. and Baltimore, D. (1993) Genes Dev., 7, 1266–1276.
- Seed, B. and Aruffo, A. (1987) Proc. Natl Acad. Sci. USA, 84, 3365-3369.
- Shu,H.B., Agranoff,A.B., Nabel,E.G., Leung,K., Duckett,C.S., Neish,A.S., Collins,T. and Nabel,G.J. (1993) *Mol. Cell. Biol.*, **13**, 6283–6289.
- Sica,A., Tan,T.-H., Rice,N., Ketzschmar,M., Ghosh,P. and Young,H.A. (1992) Proc. Natl Acad. Sci. USA, 89, 1740–1744.
- Sun,S.-C., Ganchi,P.A., Ballard,D.W. and Greene,W.C. (1993) Science, 259, 1912–1915.
- Sun,S.-C., Ganchi,P.A., Beraud,C., Ballard,D.W. and Greene,W.C. (1994) Proc. Natl Acad. Sci. USA, 91, 1346–1350.
- Tan,T.-H., Huang,G.P., Sica,A., Ghosh,P., Young,H.A., Longo,D.L. and Rice,N.R. (1992) Mol. Cell. Biol., 12, 4067–4075.
- Tewari, M., Dobrzanski, P., Mohn, K.L., Cressman, D.E., Hsu, J.-C., Bravo, R. and Taub, R. (1992) *Mol. Cell. Biol.*, **12**, 2898–2908.

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