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Activation and attenuation of transcription factor NF-kB in mouse glomerular mesangial cells in response to tumor necrosis factor-alpha, immunoglobulin G, and adenosine 3':5'cyclic monophosphate. Evidence for involvement of reactive oxygen species.

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Research Article

The transcription factor NF-kB may play an important role in the response to tissue injury and activation of cytokines. We therefore examined the regulation of NF-kB in mesangial cells. Treatment of mesangial cells with TNF-alpha increased nuclear proteins that bound to an NF-kB-specific DNA oligonucleotide. IgG aggregates also increased nuclear NF-kB demonstrating Fc-tau receptor-mediated activation of NF-kB. Treatment of a cytosolic preparation with the detergent deoxycholate also activated NF-kB. The binding characteristics were typical for NF-kB transcription factors as determined by competition experiments with NF-kB-binding wild type kB DNA oligonucleotides or mutated oligonucleotides. Furthermore, a monoclonal antibody against the p65 subunit of NF-kB prevented the binding of NF-kB to the kB oligonucleotide. To evaluate the potential role of reactive oxygen intermediates in the activation of NF-kB, we used PDTC as a scavenger and HMAP as an inhibitor of NADPH-dependent oxidase. Both PDTC and HMAP attenuated the increase in nuclear NF-kB in response to either TNF-alpha or IgG complexes. Finally, generation of superoxide anion by xanthine oxidase activated NF-kB, an effect also mitigated by PDTC. In contrast, exogenous H2O2 did not activate NF-kB. Preincubation of cells with 8 br-cAMP, forskolin, or PGE2 attenuated the increase in nuclear NF-kB in response to TNF-alpha, aggregated IgG, or superoxide anion. Our results provide support for a role of reactive oxygen intermediates as mediators for [...]



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Activation and Attenuation of Transcription Factor NF-kB in Mouse Glomerular Mesangial Cells in Response to Tumor Necrosis Factor- α , Immunoglobulin G, and Adenosine 3':5'-Cyclic Monophosphate

Evidence for Involvement of Reactive Oxygen Species

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Abstract

Introduction

The transcription factor NF-kB may play an important role in the response to tissue injury and activation of cytokines. We therefore examined the regulation of NF-kB in mesangial cells. Treatment of mesangial cells with TNF- α increased nuclear proteins that bound to an NF-kB-specific DNA oligonucleotide. IgG aggregates also increased nuclear NF-kB, demonstrating Fc- τ receptor-mediated activation of NF-kB. Treatment of a cytosolic preparation with the detergent deoxycholate also activated NF-kB. The binding characteristics were typical for NF-kB transcription factors as determined by competition experiments with NF-kBbinding wild type kB DNA oligonucleotides or mutated oligonucleotides. Furthermore, a monoclonal antibody against the p65 subunit of NF-kB prevented the binding of NF-kB to the kB oligonucleotide. To evaluate the potential role of reactive oxygen intermediates in the activation of NF-kB, we used PDTC as a scavenger and HMAP as an inhibitor of NADPH-dependent oxidase. Both PDTC and HMAP attenuated the increase in nuclear NF-kB in response to either TNF- α or IgG complexes. Finally, generation of superoxide anion by xanthine oxidase activated NF-kB, an effect also mitigated by PDTC. In contrast, exogenous H₂O₂ did not activate NF-kB. Preincubation of cells with 8 br-cAMP, forskolin, or PGE₂ attenuated the increase in nuclear NF-kB in response to TNF- α , aggregated IgG, or superoxide anion. Our results provide support for a role of reactive oxygen intermediates as mediators for activation of NF-kB in MC after stimulation with TNF- α or IgG aggregates. As an unexpected novel finding we report that cAMP can inhibit activation of NF-kB in MC. These observations may help to explain effects of TNF- α , IgG aggregates and cAMP on generation of cytokines by mesangial cells and the resulting glomerular pathophysiology. (J. Clin. Invest. 1994. 94:1629-1636). Key words: chemotactic factors • kidney • gene regulation

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Mesangial cells $(MC)^1$ are vascular smooth muscle-derived cells that play an important role in glomerular function in the kidney (1). MC also contribute to glomerular injury by generating a number of cytokines. We recently described that $TNF-\alpha$ and IgG complexes can activate expression of the macrophage colony stimulating factor CSF-1 and monocyte chemoattractant protein MCP-1. Furthermore superoxide anion activated these genes and scavengers of free radicals attenuated these effects, implicating reactive oxygen intermediates as possible mediators of gene activation.

In eucaryotes, regulation of gene transcription can be dramatically enhanced by the binding of sequence-specific DNA binding proteins to a promoter's *cis*-acting element (2, 3). Nuclear factor kB (NF-kB) is a well studied pleiotropic, multisubunit transcription factor protein originally described as binding to an immunoglobulin k enhancer site (kB) of B cells, and required for k enhancer function (4). Subsequently, a wide variety of inducible genes has been shown to contain kB binding sites in enhancer regions (5).

NF-kB resides in the cytoplasm of different cell types as an inactive complex of the subunits p50, p65, and an inhibitory subunit (IkB). The p50 and p65 subunits are the best characterized members of a larger protein family (NF-kB/Rel/Dorsal). Agents that lead to immune, inflammatory, or acute phase response activate the transcription factor by release of IkB with subsequent translocation of the p50–p65 heterodimer into the nucleus. IkB can also be dissociated from NF-kB in vitro by treatment with the detergent sodium deoxycholate (DOC) (6). As activation is independent of new protein synthesis, NF-kB has been proposed to serve as an integral factor in rapidly inducing an array of genes for signaling and defense proteins (5, 7). Cytokines, immunoglobulins, major histocompatability antigens, and their associated proteins are among those genes responsive to NF-kB activity (8).

Phosphorylation may play an important role in the activation and metabolism of NF-kB, but the contribution of various kinases to this process remains to be defined (9-13). NF-kB also appears to be influenced by reactive oxygen intermediates (ROI). ROI are commonly produced during inflammatory pro-

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^{1.} Abbreviations used in this paper: CHX, cycloheximide; DOC, sodium deoxycholate; Fsk, Forskolin; HMAP, 4'-hydroxy-3'-methoxy-acetophenone; IkB, inhibitory subunit of NF-kB; MC, mesangial cell; MCP, monocyte chemoattractant protein; NF-kB, nuclear factor kB; PDTC, pyrolidine dithiocarbamate; PKA, cAMP-dependent protein kinase; PKC, protein kinase C; ROI, reactive oxygen intermediate; XO, xanthine oxidase-hypoxanthine.

cesses. In mesangial cells TNF- α (14) and immune complexes (15) induce the production of ROI. Furthermore, MC contain an inducible, membrane bound NADPH-dependent oxidase capable of H₂O₂ and O₂⁻ generation (16). Oxygen radicals, in turn, have recently been implicated as activators of NF-kB and may be a common mechanism of action for the diverse agents capable of NF-kB activation (17–20).

Macrophage colony stimulating factor (CSF-1) and monocyte chemoattractant protein (MCP-1 also known as MCAF) are two monocyte-specific inducible cytokines that contain a kB site upstream from their promoters (21, 22). CSF-1 and MCP-1 genes can be rapidly induced by a number of agents including lipopolysaccharide, PDGF, TNF- α , IL-1. IFN- τ , and occupancy of Fc receptors by aggregated IgG (22-36). We also have shown that cAMP and agents increasing cAMP generation such as forskolin and PGE₂ attenuate the increase in mRNA for CSF-1 and MCP-1 in response to TNF- α or aggregated IgG (37). Recently we demonstrated that TNF- α and aggregated IgG can stimulate NADPH-dependent generation of free radicals which may serve as intracellular signals for the induction of CSF-1 and MCP-1 in glomerular mesangial cells (38). Furthermore, radical scavengers including N-acetyl-L-cysteine (NAC) and pyrolidine dithiocarbamate (PDTC) were able to attenuate the TNF- α and IgG induced expression of MCP-1 (38).

We now report activation of NF-kB by treatment of MC with TNF- α and, for the first time, with Fc- γ receptor occupancy by aggregated IgG. 8 br-cAMP or agents which stimulate cAMP such as forskolin or PGE₂ attenuate the induction of NF-kB by TNF- α or aggregated IgG. These results extend our previous studies on the transcriptional, mRNA, and protein levels of CSF-1 and MCP-1 in activated MC (36–38). Through the use of the NADPH oxidase inhibitor 4'-hydroxy-3'-methoxy-acetophenone (HMAP), the free radical scavenger PDTC, and the superoxide anion generating xanthine oxidase–hypoxanthine system, we provide evidence that implicate ROI generation as a common pathway for activation of NF-kB by TNF- α and aggregated IgG in MC.

Methods

BSA, forskolin, PGE₂, PDTC, and 8 bromo-cAMP were from Sigma Chemical Co. (St. Louis, MO). DME, penicillin (0.66 mg/ml)-streptomycin (60 μ g/ml), and FCS were purchased from Gibco BRL (Grand Island, NY). HMAP was purchased from Aldrich Chemical Co. (Milwaukee, WI). Mouse recombinant TNF- α , acrylamide, cycloheximide (CHX), and multiprime DNA labeling systems were obtained from Boehringer Mannheim (Indianapolis, IN). [α -³²P]dCTP was purchased from New England Nuclear (Boston, MA). NF-kB p65 subunit specific monoclonal antibody was purchased from PharMingen (San Diego, CA). Mouse IgG or F(ab')₂ fragment (Cappel Organon Teknika Corp., Durham, NC) was aggregated by heating to 63°C for 30 min. Insoluble aggregates were removed by centrifugation at 1,000 rpm in a microfuge for 1 min and the supernate served as aggregated IgG or F(ab')₂ fragments, respectively (36). All other chemicals were from Sigma Chemical Co., unless otherwise specified.

Culture of mouse MC. Mouse MC transformed with nonreplicating, noncapsid forming SV-40 virus (strain Rh 911) were kindly provided by Dr. E. Neilson, University of Pennsylvania. These cells have been previously characterized (39). In previous experiments we had established that the response to TNF α and aggregated IgG was comparable in the transformed MC and primary MC (38). The MC were maintained in DME medium with 1% Penicillin–Streptomycin and 2.5% FCS in an atmosphere of $95\% O_2$, $5\% CO_2$. Medium was changed twice weekly. Cells were passaged weekly by harvesting with 0.25% trypsin, 1 mM EDTA solution.

Preparation of nuclear extracts. For 16 h before the experiments cells were kept in medium with the omission of FCS. Incubation media, ±experimental agents, were also without FCS. Preincubation with PDTC (0.1 mM), HMAP (100 µg/ml), CHX (15 µg/ml), 8br-cAMP (1 mM), forskolin (Fsk; 10⁻⁵ M) or PGE₂ (10⁻⁶ m) for 30 min or 1 h as specified. Cells were treated with TNF- α (100 U/ml) or aggregated IgG (200 µg/ml), XO (xanthine oxidase + hypoxanthine at 10 mU/ ml and 0.5 mM, respectively), or hydrogen peroxide (50 mM) for an additional 1 h at 37°C.

Nuclear proteins were isolated by the method of Schriber et al. (40). In brief, confluent MC (5 \times 10⁶ cells) were treated with experimental agents for the times indicated, the media were then quickly aspirated and cells washed gently on ice with 10 ml ice-cold Tris-buffered saline, pH 7.9. Cells were scraped in 1 ml of cold hypotonic buffer A (10 mM Hepes, pH 7.9; 10 mM KCl, 0.1 mM EDTA; 0.1 mM EGTA; 1 mM DTT; 0.5 mM PMSF) + 0.6% NP-40, allowed to swell on ice for 15 min and vortexed vigorously for lysis. After a 30-s centrifugation in a microfuge, the pellet was then washed with an additional 1 ml cold buffer and respun. The resulting nuclear pellet was resuspended in 100 µl ice-cold buffer C (20 mM Hepes pH 7.9; 0.4 M NaCl; 1 mM EDTA; 1 mM EGTA; 1 mM DTT; 1 mM PMSF) and placed in an Eppendorf thermomixer at setting 10 for 20 min at 4°C. The nuclear extract was centrifuged for 10 min at 11,500 g at 4°C; the supernatant collected, divided into aliquots and stored at -70°C. Protein concentration was determined by the Bio-Rad Protein Assay. 20 μ g of this extract, typically $1-2 \mu l$, was used for the bandshift assay.

For DOC pretreatment, cytosolic fraction $(25 \ \mu g)$ is incubated for 10 min at room temperature with DOC (0.5% wt/vol). DOC is then sequestered by addition of NP-40 (1%) immediately before EMSA binding reaction (see EMSA in Methods).

Electrophoresis mobility shift assay (EMSA). The kB DNA sequence of the immunoglobulin gene (6) was used for the EMSA. Synthetic double stranded oligonucleotides of the following sequences were end labeled with $[^{32}P]$:

kB:	5 'TCGACAGAGGGACTTTCCGAGAGGC
	3' GTCTCCCTGAAAGGCTCTCCGAGCT
mutated kB:	5 'TCGACAGAATTCACTTTCCGAGAGGC
	3' GTCT <u>TAAG</u> TGAAAGGCTCTCCGAGCT
TRE:	5'GATGCAGCTTGATGAGTCAGCCGGATCA 3' GTCGAACTACTCAGTCGGCCTAGTCTAG

The altered sequence of the mutated DNA sequence is underlined.

The DNA binding reaction was performed on ice for 20 min in a volume of 20 μ l, unless otherwise specified. The reaction mixture contained 20 μ g of nuclear extract, 50 μ g/ml of poly dI:dC, 10 mM Tris pH 7.5, 100 mM NaCl, 1 mM EDTA, 1 mM DTT, 1 mg/ml BSA, 10% glycerol and 25,000 cpm (³²P) end labeled oligonucleotide. Following incubation, the samples were loaded onto a 5% native polyacrylamide gel (acrylamide/bisacrylamide at 40:1 in 0.3× TBE) while running at 100 V. After loading, the voltage was increased to 150 V for 2 h. The gel was then dried and subjected to autoradiography for analysis.

For binding exclusion assays, anti–NF-kB p65 is added either before or after labeled kB oligonucleotide addition, as noted, for 20 min incubation at 37° C.

PKA induction of NF-kB. Nuclear protein extracts $(20 \ \mu g)$ or cytosolic fractions $(50 \ \mu g)$ were incubated in the presence or absence of 1 mM MgCl₂, 100 μ M ATP or ATP gamma-S, a nonhydrolizable ATP analogue, and 0.2 μ g catalytic subunit of cAMP-dependent protein kinase (PKA). The DNA binding reaction, as described above in EMSA, was performed immediately after the PKA incubation.

Results

kB Sequence binds NF-kB-like protein in MC. To detect kB binding activity in MC, we performed electrophoretic mobility



Preincubation

tions of Poly dI-dC as a nonspecific competitor on binding of labeled kB oligonucleotide sequence to nuclear extracts from MC preincubated with TNF- α (100 U/ml). Brackets indicate position of NF-kB bands. (B) Binding of nuclear extract from TNF- α stimulated MC to labeled mutant kB sequence (lane 1; see text) or to labeled control kB sequence (remaining lanes). Competition of binding was performed with increasing concentrations of unlabeled kB oligonucleotide, unlabeled kB-mutant oligonucleotide, or unlabeled TRE oligonucleotide. Concentration of Poly dI-dC was 50 μ g/ml. (C) Binding of cytosolic fractions (25 μ g) of MC lysate to labeled kB oligonucleotide without (-) or with (+) DOC preincubation at room temperature for 10 min (results from a duplicate experiment). (D) Binding of nuclear extract from TNF- α stimulated MC (5 μ g) without or with addition of monoclonal antibody against the p65 subunit of NF-kB (anti-NF-kB). Nuclear extracts were preincubated for 20 min at 37°C without (TNF- α alone) or with $(TNF-\alpha + anti-NF-kB)$ increasing concentrations of antibody prior to addition of labeled kB oligonucleotide to the binding reaction. Increasing concentrations of the antibody causes exclusion of the NF-kB bands. (Preincubation, lanes 2-8). When binding of nuclear extract and labeled kB sequence were performed prior to addition of antibody the NF-kB bands were not excluded (Post-incubation, lanes 9 and 10).

Figure 1. EMSA analysis for NF-kB activity in MC. (A) Effects of increasing concentra-

shift assays (EMSA) using a 26-bp, end labeled oligonucleotide recognizing NF-kB of the immunoglobulin gene (kB) as described by Zabel et al. (6). As shown in Fig. 1 A, nuclear protein extracts from MC stimulated with TNF- α caused band retardation of the labeled kB DNA sequence. In preliminary experiments, using from 5 to 30 μ g of nuclear protein extracts, we established that optimal concentrations for the EMSA were present at 15–25 μ g of nuclear extract. Increasing concentrations of poly dI-dC as nonspecific competitor displaced multiple bands considered to be nonspecific. Poly dI-dC did not displace

cytosolic fractions

binding of bands with an NF-kB like motility even at 250 μ g/ml (Fig. 1 *A*). There are several bands on the EMSA which most likely correspond to different NF-kB like factor subunit combinations. These NF-kB like bands could effectively be competed out with increasing concentrations of unlabeled NF-kB binding DNA consensus sequence (Fig. 1 *B*). In contrast, the mutated kB sequence was far less efficient (~ 20-fold, as determined by densitometry) in binding nuclear proteins or in displacing labeled NF-kB binding DNA sequences indicating kB-specific DNA binding of the nuclear protein extracts. Fur-

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thermore, an oligonucleotide corresponding to the TPA regulatory element (TRE) recognition sequence was an ineffective competitor even at 250-fold excess (Fig. 1 B).

DOC has been shown to release the heterodimer NF-kB from IkB in vitro (41). EMSA on the cytosolic fraction of MC with or without pretreatment with DOC showed a marked increase in NF-kB binding after DOC pretreatment indicating that MC cytosol contains inactive, inducible NF-kB (Fig. 1 C).

NF-kB-like factor binding could be progressively excluded by preincubating with increasing concentrations of NF-kB monoclonal antibody (Fig. 1 D). Thus the antibody prevents binding of the NF-kB to the kB sequence. Conversely no decrease in labeled kB oligonucleotide binding was observed when anti-NF-kB was added after the normal EMSA binding reaction. This would indicate that the epitope on the NF-kB is no longer accessible to the antibody once NF-kB is bound to the DNA element.

TNF- α or aggregated IgG activate NF-kB in MC: effects attenuated by cAMP. Previously we have shown that exposure of MC to TNF- α or aggregated IgG results in increased expression of CSF-1 and MCP-1 with concurrent increases in the levels of their respective mRNA (36, 37). Conversely, cAMP downregulates the expression of these genes in MC (37). We therefore examined if TNF- α , aggregated IgG, cAMP, and cAMP mimetics such as forskolin (Fsk) or PGE₂ could influence the activation and translocation of NF-kB. As shown in Fig. 2 A, NF-kB binding activity of MC nuclear extracts increased when cells had been treated with TNF- α or aggregated IgG. The $F(ab')_2$ fragment of IgG only resulted in a minor band-shift. This may, at least in part, be due to uncleaved IgG in the $F(ab')_2$ preparation. This is consistent with activation of NF-kB by Fc gamma receptor occupancy by IgG. The effect of TNF- α and aggregated IgG were attenuated when cells were preincubated with cAMP, forskolin or PGE₂. In contrast, pretreatment of MC with these experimental agents had no effect on the binding of nuclear extracts to the TPA regulatory element (TRE, a member of the AP-1 family; Fig. 2 B). Specificity of nuclear protein binding to the TRE oligonucleotide had been established in preliminary experiments using labeled mutated TRE oligonucleotide and displacement by unlabeled TRE and mutated TRE oligonucleotides (results not shown). This argues against a nonspecific effect of pretreatment of MC with TNF- α , IgG, cAMP, forskolin, or PGE₂ on EMSA.

Activation of NF-kB also occurred after treatment with cycloheximide (CHX), indicating that NF-kB can be activated independent from de novo protein synthesis as previously reported by Sen and Baltimore (4) (Fig. 2 C). The effect of cycloheximide has been related to the inhibitory subunit of NF-kB (42).

In vitro treatment with protein kinase A does not effect NFkB EMSA. As cAMP decreased activation of NF-kB in response to TNF- α , we examined whether this effect could be mimicked by PKA treatment of cell extracts. When nuclear or cytosolic extracts from MC pre-exposed to TNF- α were incubated with MgCl₂, \pm ATP or nonhydrolizable ATP gamma-S \pm catalytic subunit of PKA, no change in gel shift occurred (Fig. 3). Comparable results were obtained with extracts from cells that had not been exposed to TNF- α (results not shown). Similar results were obtained when ATP concentrations were raised to 1 and 10 mM ATP (results not shown). Thus in vitro treatment with PKA did not induce activation of cytosolic NF-kB or



Figure 2. Effects of TNF- α (100 U/ml), aggr. IgG (200 μ g/ml) or F(ab')₂ (200 μ g/ml) alone or in combination with 8br-cAMP (1 mM), forskolin (Fsk; 10⁻⁵/M) or PGE₂ (10⁻⁶/M) on EMSA for nuclear protein extracts binding to labeled kb oligonucleotide (Fig. 2 *A*) or labeled TRE oligonucleotide (Fig. 2 *B*). Cycloheximide (CHX; 15 μ g/ μ l) also activated NF-kB (Fig. 2 *C*). MC were preincubated for 30 min with 8br-cAMP, Fsk or PGE₂ prior to a 1-h co-incubation±TNF- α or aggregated IgG before extraction of nuclear proteins and EMSA. Comparable results were obtained in 5 sets of experiments for 8br-cAMP, 3 sets for forskolin and 4 sets for PGE₂.

influence binding of the nuclear factor to the labeled kB oligonucleotide.

Generation of superoxide activates induction of NF-kB. It has been proposed that NF-kB can be activated by reactive oxygen intermediates and that activation can be suppressed by thiol containing radical scavengers such as NAC and PDTC (17-20). We recently provided evidence that reactive oxygen species may be involved in raising mRNA levels for MCP-1 and CSF-1 (38) consistent with the hypothesis that free radicals



Figure 3. Effects of preincubation of nuclear or cytosolic extracts with protein kinase A. Cells were stimulated with TNF- α prior to extraction. 20 μ g nuclear protein or 50 μ g cytosolic extracts were preincubated with MgCl₂ (1 mM)±ATP (100 μ M) or ATP gamma-S (100 μ M)±PKA (0.2 μ g catalytic subunit) for 10 min at 30°C prior to binding to labeled kB sequence and EMSA.

can serve as mediators in gene activation (43). We now examined if free radicals could also activate NF-kB in MC. As a thiol agent we used PDTC (20). Pretreatment of MC with PDTC partially but consistently inhibited the activation of NFkB in response to either TNF- α or aggregated IgG (Fig. 4 A). These findings are consistent with free radicals serving as intermediates in NF-kB activation.

A membrane bound NADPH-dependent oxidase system has been implicated for the generation of oxygen free radicals in stimulated MC (16, 38). HMAP, also known as apocynin, a reported inhibitor of NADPH-dependent oxidase (44), can decrease superoxide generation by MC stimulated with either TNF- α or aggregated IgG and concurrently decrease expression of mRNA for CSF-1 and MCP-1 (38). We therefore used HMAP to see if this agent would also decrease activation of NF-kB by either TNF- α or aggregated IgG. Results with HMAP were similar to those with PDTC (Fig. 4 B), i.e., HMAP attenuated the activation of NF-kB by TNF- α or IgG complexes. Again, EMSA using labeled TRE sequence were unaffected by TNF- α or aggregated IgG. In contrast to the attenuation of the NF-kB binding in the gel shifts after pretreatment in the presence of PDTC, PDTC enhances the EMSA for TRE (results not shown). Enhancement of AP-1 binding by PDTC has recently been described (45).

To examine whether generation of O_2^- was also capable of activating NF-kB, we used generation of exogenous O_2^- by the addition of xanthine oxidase and hypoxanthine (14) to the incubation media. Heat inactivated xanthine oxidase (by boiling the

enzyme for 3 min; iXO) served as control. As shown in Fig. 5 A, generation of O_2^- activated NF-kB in MC, whereas treatment with heat inactivated XO had no effect. Furthermore the radical scavenger PDTC, as well as 8br-cAMP, also attenuated the activation of NF-kB by O_2^- . In contrast, H_2O_2 demonstrated no discernible effect on NF-kB in the EMSA (Fig. 5 B). These results are in accord with our earlier studies where generation of O_2^- , but not H_2O_2 , stimulated mRNA levels of CSF-1 and MCP-1 (38).

Discussion

Our present results demonstrate a parallelism between the activation and attenuation of NF-kB activity in MC and our previous results for expression of mRNA for CSF-1 and MCP-1. Enhanced kB binding activity is produced in response to treatment with TNF- α , aggregated IgG as well as by generation of O_2^- by the xanthine oxidase-hypoxanthine system. This kB binding activity appears indistinguishable from native NF-kB in that: (a) binding could be effectively competed out only with unlabeled NF-kB consensus sequence, but not with mutated kB sequence or the TRE consensus sequence (Fig. 1 B); (b) dissociation of cytosolic fractions of MC by DOC can release NF-kB from its cryptic form (Fig. 1 C), (41); (c) NF-kB-like binding activity can be induced in EMSA by pretreatment with CHX (Fig. 2 C), (46); (d) kB binding activity is excluded when monoclonal antibodies to p65 subunits are added prior to the binding reaction (Fig. 1 D).

NF-kB has been implicated as a signal for cell distress in response to many adverse conditions, in which it can rapidly activate expression of inflammatory, immune, and acute phase response genes (5, 7). Two such genes that contain kB sequences upstream from their promoters and may fall under the regulation of NF-kB include the monocyte specific cytokines CSF-1 and MCP-1 (21, 22). MCP-1 and CSF-1 have been implicated in instituting and maintaining influx, proliferation, activation, and survival of monocyte-macrophages in immune-mediated diseases such as glomerulonephritis (36-38, 47). We recently demonstrated that transcription, as well as levels of mRNA and protein for CSF-1 and MCP-1 are increased by TNF- α or aggregated IgG and attenuated by cAMP or cAMP mimetics such as forskolin or PGE₂ (37). Our previous results thus correlate with the increased kB binding activity of MC stimulated by TNF- α or aggregated IgG and the diminished gel shift after cAMP pretreatment demonstrated in our present study (Fig. 2A).

In vitro highly purified NF-kB can be activated by phosphorylation, presumably at the IkB subunit, by cyclic AMP-dependent protein kinase (PKA) or protein kinase C (PKC) (9). Attempts were less successful with crude cytosolic extracts (9), with the exception of a study by Shirakawa and Mizel (48). Using 70Z/3 murine pre-B cells and a human natural killerlike cell line YT, these authors were able to demonstrate activation of NF-kB catalyzed by PKA or PKC phosphorylation. In our studies treatment of cytosolic or nuclear preparations with PKA neither activated nor inactivated NF-kB in MC (Fig. 3). Similar findings have been reported in the T cell lymphoma Jurkat cell line (10), or the monocyte cell line HL-60 (29). PKA, unlike PKC, was also unable to phosphorylate highly purified IkB in vitro (9). Surprisingly, pretreatment of cells



with cAMP or cAMP mimetics decreased the amount of nuclear NF-kB binding activity as determined by EMSA. This inhibitory effect of cAMP on nuclear NF-kB is apparent after stimulation of cells with TNF- α , IgG complexes, or generation of superoxide anion. Thus in intact MC cAMP inhibits activation of NF-



Figure 5. (A) Generation of superoxide anion by xanthine oxidase plus hypoxanthine (XO; 10 mU/ml and 0.5 mM, respectively) activates NFkB. Heat inactivated xanthine oxidase (iXO) does not activate NF-kB. PDTC and cAMP inhibit activation of NF-kB in response to superoxide generation by xanthine oxidase (XO + PDTC and XO + 8br-cAMP). PDTC or 8br-cAMP by themselves had no effects. H₂O₂ (50 μ M) does not activate NF-kB (Fig. 5 *B*). Comparable results were obtained in three sets of experiments. MC were preincubated for 1 h with PDTC or for 30 min with 8br-cAMP followed by a 1-h incubation±XO or H₂O₂. Nuclear proteins were extracted and bound to labeled kB as per Methods.

Figure 4. Effects of (A) PDTC (0.1 mM) or (B) HMAP (100 μ g/ml) on NF-kB activation by TNF- α or aggr. IgG as demonstrated by EMSA. Comparable results were obtained in 9 sets of experiments for PDTC and 4 sets of experiments for HMAP. Preincubation of MC with PDTC or HMAP was for 1 h at 37°C before a 1-h incubation±TNF- α or aggregated IgG. Nuclear proteins were extracted and allowed to bind with labeled kB oligonucleotide as described in Methods.

kB. Cyclic AMP could interfere with activation of NF-kB by TNF- α or IgG at multiple steps, e.g., (*a*) with signals leading to NF-kB activation; (*b*) with NF-kB activation itself; (*c*) with transfer of NF-kB to the nucleus; or (*d*) with NF-kB binding to its DNA-recognition site. The fact that cAMP also decreased activation of NF-kB in response to superoxide anion would indicate that cAMP can interfere with steps downstream of O_2^- generation. Our experiments using PKA catalytic subunit in broken cell preparations only indicate that in vitro treatment of cytosolic or nuclear fractions have no influence on NF-kB dissociation, or binding to the NF-kB specific oligonucleotide. Future experiments will have to examine the mechanism for the negative regulation of NF-kB by cAMP.

All agents examined induced the release of reactive oxygen intermediates (ROI). These intermediates have been implicated as second messengers in the induction of NF-kB (17-20). The ROI could be generated by NADPH oxidase, a membrane bound enzyme (49) found, in addition to phagocytic cell lines, in cells of mesenchymal origin, including MC (16, 38). This study shows that scavengers of ROI, such as PDTC, can attenuate activation of NF-kB in MC. HMAP, an inhibitor of ROI generation by NADPH-dependent oxidase (38, 44) also reduces activation of NF-kB, arguing in favor of ROI playing a role in NFkB activation. Generation of superoxide anion, but not H_2O_2 , activates NF-kB providing an additional argument in favor of ROI as second messengers.

We find it intriguing to speculate on the correlation between regulation of mRNA levels for example, MCP-1 and CSF-1 in MC and activation of NF-kB. Thus cAMP decreases mRNA levels and EMSA for NF-kB in MC stimulated with TNF- α or IgG. Scavengers of free radicals or inhibition of their formation by HMAP also shows concordant effects on mRNA levels and NF-kB gel shifts. Generation of superoxide anion activates NFkB and increases mRNA levels for MCP-1 and CSF-1. While this correlation is highly intriguing it does not establish causality, which will require future experiments using reporter gene assays.

Overall our results support the proposed role for ROI in NF-

kB activation and the resulting transcription of various genes including MCP-1 and CSF-1 involved in acute tissue injury. The attenuating effects of cAMP and radical scavengers on this process may indicate potential avenues for therapeutic interventions during glomerular injury, for example.

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