



PKC- and ERK-dependent activation of I κ B kinase by lipopolysaccharide in macrophages: enhancement by P2Y receptor-mediated CaMK activation

¹Bing-C. Chen & ^{*,1}Wan-W. Lin

¹Department of Pharmacology, College of Medicine, National Taiwan University, Taipei, Taiwan

1 Although accumulating studies have identified I κ B kinase (IKK) to be essential for controlling NF- κ B activity in response to several cytokines, the upstream kinases that control IKK activity are still not completely known. We have previously reported that G protein-coupled P2Y₆ receptor activation by UTP potentiates lipopolysaccharide (LPS)-induced I κ B phosphorylation and degradation, and NF- κ B activation in J774 macrophages. In this study, we investigated the upstream kinases for IKK activation by UTP and LPS.

2 In murine J774 macrophages, LPS-induced NF- κ B activation was inhibited by the presence of PDTC, D609, Ro 31-8220, PD 098059 and SB 203580.

3 Accompanying NF- κ B activation, LPS induced I κ B degradation and IKK activation were reduced by PDTC, D609, Ro 31-8220 and PD 098059, but not by SB 203580.

4 Although UTP itself slightly induced IKK activation, this response was synergistic with LPS. BAPTA/AM and KN-93 (a calcium/calmodulin-dependent protein kinase (CaMK) inhibitor) attenuated UTP- but not LPS-stimulated IKK activity. Synergistic IKK activation between LPS and thapsigargin was further demonstrated in peritoneal macrophages.

5 LPS and UTP co-stimulation additively increased p65 NF- κ B phosphorylation. *In vitro* kinase assays revealed that LPS and UTP induced extracellular signal-regulated protein kinase (ERK) and p38 mitogen-activated protein kinase activation were respectively inhibited by PD098059 and SB 203580.

6 Taken together, we demonstrate that Gq protein-coupled P2Y₆ receptor activation can potentiate LPS-stimulated IKK activity. While PKC and ERK participate in IKK activation by LPS and UTP, the phosphatidylinositol-3-phospholipase C-dependent activation of CaMK plays a major role in UTP potentiation of the LPS response.

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Abbreviations: CaMK, calcium/calmodulin-dependent protein kinase; COX-2, cyclooxygenase-2; DMEM, Dulbecco's modified Eagle's medium; ECL, enhanced chemiluminescence; EMSA, electrophoretic mobility shift assay; ERK, extracellular signal-regulated protein kinase; IKK, I κ B kinase; IL-6, interleukin-6; iNOS, inducible nitric oxide synthase; LPS, lipopolysaccharide; MAPK, mitogen-activated protein kinase; MBP, myelin basic protein; MEK, mitogen-activated protein kinase kinase; NIK, NF- κ B-inducing kinase; PBS, phosphate-buffered saline; PC, phosphatidylcholine; PDTC, pyrrolidine dithiocarbamate; PGE₂, prostaglandin E₂; PI, phosphatidylinositol; PKC, protein kinase C; PLC, phospholipase C; PMA, phorbol 12-myristate 13-acetate; TG, thapsigargin; TNF α , tumor necrosis factor- α

Introduction

The transcription factor, NF- κ B, plays a key role in the transcriptional regulation of many proteins involved in chronic inflammatory diseases (Barnes & Karin, 1997). NF- κ B is a dimer of transcription factors primarily consisting of p65 (RelA) or p50; p65 is the transcriptional activation component of the most common form of the NF- κ B heterodimer (Baldwin, 1996). In the resting state, this dimer is anchored by I κ B α and I κ B β , which function to retain NF- κ B in the cytosol (Thompson *et al.*, 1995). Recently, two closely related I κ B kinases (IKKs), IKK α and IKK β , were identified which directly phosphorylate I κ B α at Ser³² and Ser³⁶, and phosphorylate I κ B β at Ser¹⁹ and Ser²³ (DiDonato

et al., 1997; Maniatis, 1997; Woronicz *et al.*, 1997). These phosphorylations lead to ubiquitination of I κ Bs at specific lysine residues, and their degradation by the 26S proteasome (Chen *et al.*, 1995). This process releases active NF- κ B, which then translocates to the nucleus, binds to specific DNA enhancer sequences (κ B binding sites), and activates gene transcription. Based on approaches using transient transfection overexpression, the best-characterized upstream kinase candidate involved in the activation of the IKK complex in response to specific stimuli through phosphorylation of the two IKK subunits has been identified as NF- κ B-inducing kinase (NIK) (Malinin *et al.*, 1997). All NF- κ B activation either by cytokines, such as interleukin-1 and tumour necrosis factor- α (TNF α), or by bacterial endotoxin lipopolysaccharide (LPS) requires NIK activation (Israel, 2000).

*Author for correspondence; E-mail: wwl@ha.mc.ntu.edu.tw

Transcriptional activity of NF- κ B can be regulated by mechanisms other than cytosolic degradation of I κ B. Several reports have demonstrated that the DNA binding (Hayashi *et al.*, 1993) and transactivating capacity (Zhong *et al.*, 1997; Wang & Baldwin, 1998) of NF- κ B are up-regulated by inducible phosphorylation of p65. In this context, protein kinases, such as IKKs (Sakurai *et al.*, 1999), protein kinase A (PKA) (Zhong *et al.*, 1998), p38 mitogen-activated protein kinase (MAPK) (Vanden Berghe *et al.*, 1998; Jefferies & O'Neill, 2000), extracellular signal-regulated protein kinase (ERK) (Vanden Berghe *et al.*, 1998; Jefferies & O'Neill, 2000), and PKC (Lozano *et al.*, 1994) have been shown to phosphorylate p65, for example at Ser²⁷⁶ by PKA or at Ser⁵³⁶ by IKKs. This p65 phosphorylation in turn increases the transcriptional activity of NF- κ B, and represents another mechanism, independent of I κ B degradation, for enhancing NF- κ B activation.

Although there has been a tremendous increase in our understanding of the mechanisms leading to NF- κ B activation during the past 3 years, it is still unclear whether other kinases, with the exception of NIK, mediate the regulation of IKK complex activity. One of the reasons this unanswered question has been raised is that recent data have implicated mitogen-activated protein kinase/ERK kinase kinase 1 (MEKK1) (Lee *et al.*, 1997), PKC, ERK, and p38 MAPK in NF- κ B activation. With respect to PKC, PKC ζ was shown to phosphorylate I κ B α *in vitro* (Diaz Meco *et al.*, 1994), and transfection of a dominant negative mutant of PKC ζ severely impairs NF- κ B activation by sphingomyelinase (Lozano *et al.*, 1994). *In vivo* overexpression studies also demonstrated PKC ϵ and PKC α to be inducers of NF- κ B activation (Genot *et al.*, 1995; Trushin *et al.*, 1999). Some studies, including ours, have also demonstrated that p38 MAPK is a crucial signal kinase mediating LPS-elicited NF- κ B activation (Chen *et al.*, 1999; Nick *et al.*, 1999). At present, the roles of PKC, ERK, and p38 MAPK in LPS-induced IKK activity have yet to be investigated.

Extracellular nucleotides such as ATP and UTP exert diverse effects on cellular function by acting on P₂ receptors. In murine J774 macrophages, we have demonstrated that the expressed P2Y₆ receptors mediate UTP and UDP actions in increasing phosphoinositide (PI) turnover, PKC activation, and intracellular Ca²⁺ ([Ca²⁺]_i) mobilization (Lin & Chen, 1997). Subsequently, we showed the potentiation effects of UTP on LPS-elicited inducible nitric oxide synthase (iNOS), cyclooxygenase-2 (COX-2), and interleukin-6 (IL-6) gene induction. UTP potentiating effects are ascribed to a signalling pathway controlled by a calcium/calmodulin-dependent protein kinase (CaMK), which involves the enhancement of I κ B phosphorylation and degradation as well as NF- κ B activation (Chen *et al.*, 1998; Chen & Lin, 1999; 2000). Apart from these findings, the molecular mechanisms by which UTP potentiates I κ B phosphorylation are still unclear. Thus, in this study, we address the effects of UTP on IKK activity and the roles of CaMK, PKC, ERK, and p38 MAPK in LPS plus UTP-induced NF- κ B activation.

Methods

Cell culture

The murine J774 macrophages obtained from American Type Culture Collection were cultured in Dulbecco's modified

Eagle's medium (DMEM) containing 10% foetal bovine serum and antibiotics (100 U ml⁻¹ of penicillin and 100 μ g ml⁻¹ streptomycin). The peritoneal macrophages were prepared from Balb/c mice which had been i.p. injected with 1.5 ml of 3% thioglycolate 3 days before macrophage isolation. The peritoneal cavities were flushed with ice-cold 0.9% NaCl to remove the elicited peritoneal macrophages which were then cultured at 37°C in RPMI 1640, supplemented with 10% foetal bovine serum and antibiotics (100 U ml⁻¹ penicillin and 100 μ g ml⁻¹ streptomycin). Cells were cultured at 37°C in a humidified atmosphere of 95% air/5% CO₂.

Immunoblot analysis of I κ B α and I κ B β

To quantify I κ B α and I κ B β proteins in cells following LPS treatment for different periods, cells were washed twice in ice-cold phosphate-buffered saline (PBS), and then solubilized in buffer containing (mM): Tris-HCl 20 (pH 7.5), EGTA 0.5, EDTA 2, dithiothreitol 2, p-methylsulphonyl fluoride 0.5, and 10 μ g ml⁻¹ leupeptin. Samples of equal amounts of protein (100 μ g) were subjected to SDS-PAGE on 12% polyacrylamide gels, then transferred onto a nitrocellulose membrane, which was then incubated in TBST buffer (150 mM NaCl, 20 mM Tris-HCl, 0.02% Tween 20, pH 7.4) containing 1% nonfat milk. The I κ B α or I κ B β band was visualized by immunoblotting with a specific antibody. Immunoreactivity was detected by enhanced chemiluminescence (ECL) following the manufacturer's instructions. All the protein levels were expressed as percentages of control group.

Immunoprecipitation and protein kinase assays

Cells were washed twice with ice-cold PBS, lysed in 1 ml of lysis buffer containing: 20 mM Tris, pH 7.5, 1 mM MgCl₂, 125 mM NaCl, 1% Triton X-100, 1 mM p-methylsulphonyl fluoride, 10 μ g ml⁻¹ leupeptin, 10 μ g ml⁻¹ aprotinin, 25 mM β -glycerophosphate, 50 mM NaF, and 100 mM sodium orthovanadate, and centrifuged. The supernatant was then immunoprecipitated with respective polyclonal antibodies against IKK α , IKK β , ERK2, or p38 MAPK in the presence of A/G-agarose beads overnight. The beads were washed three times with lysis buffer and two times with kinase buffer (20 mM HEPES, pH 7.4, 20 mM MgCl₂, 2 mM dithiothreitol). Then the beads were equally divided into two parts respectively for kinase assay and immunoblotting. The kinase reactions were performed by incubating immunoprecipitated beads with 20 μ l of kinase buffer supplemented with 20 μ M ATP and 3 μ Ci of [γ -³²P]ATP at 30°C for 30 min. For IKK assay, both IKK α and IKK β antibodies were used to immunoprecipitate both kinases, and 2.5 μ g of bacterially expressed GST-I κ B α (amino acids 5–55), GST-p65 (amino acids 354–551), or GST-p65 S536A (Ser⁵³⁶ substituted by Ala⁵³⁶) was added as a substrate. For ERK and p38 MAPK assays, 50 μ g ml⁻¹ of myelin basic protein (MBP) was added as a substrate. The reaction mixtures were analysed by SDS-PAGE followed by autoradiography. All the kinase activity were expressed as percentages of the control groups. The precipitated protein levels of IKK α , ERK or p38 MAPK were assayed by immunoblotting as an internal control.

In vivo p65 phosphorylation

J774 cells were starved in medium without serum for 24 h. Then the medium was removed, replaced with phosphate-free DMEM containing 0.1 mCi ml^{-1} of [^{32}P] orthophosphate, and incubated overnight. Cells were treated with LPS, UTP, or both for different periods, followed by cell harvest in 1 ml lysis buffer and centrifugation. The supernatant was then immunoprecipitated with p65 polyclonal antibody overnight. The immunoprecipitates were then analysed by SDS-PAGE followed by autoradiography.

Preparation of nuclear extracts and electrophoretic mobility shift assays (EMSA)

Nuclear extracts were prepared as described previously (Chen *et al.*, 1998). To detect NF- κ B activation using the EMSA method, binding reaction mixtures (15 μl) contained 0.25 μg of poly (dI-dC) and 20,000 d.p.m. of ^{32}P -labelled DNA probe in binding buffer consisting of (mM): Tris-HCl 10, pH 7.5, EDTA 1, 4% Ficoll, dithiothreitol 1, and KCl 75. The binding reaction was initiated by the addition of cell extracts, and it was allowed to continue for 30 min. Samples were analysed on native 5% polyacrylamide gels. For super-shift experiments, 4 μg of anti-p65 or anti-p50 antibody was mixed with the nuclear extract proteins.

CaMK activity assay

Confluent cells on 35 mm dishes were washed with physiological saline solution (PSS, composition in mM): NaCl 118, KCl 4.7, CaCl_2 1.8, MgCl_2 1.2, KH_2PO_4 1.2, glucose 118, and HEPES 20, pH 7.4, and then the indicated drugs were added. Cells were collected on ice and disrupted by sonication at 4°C in lysis buffer. Each homogenate was immediately used for CaMK assay (Hanson & Schulman, 1992). Reaction assay mixture contained (final concentration in 50 μl): 50 mM PIPES (pH 7.0), 10 mM MgCl_2 , 0.1 mg ml^{-1} BSA, 0.3 mM CaCl_2 , 20 μM [γ - ^{32}P]ATP, 0.3 μM calmodulin, and 15 μM autocamide-3. Reactions were carried out for 30 min in 30°C , and incorporation of ^{32}P into autocamide-3 was determined. Phosphorylation of autocamide-3 in the absence of Ca^{2+} (by removing 0.3 mM CaCl_2 and addition of 0.5 mM EDTA) and calmodulin relative to its phosphorylation in the presence of Ca^{2+} and calmodulin is defined as the autonomous activity.

Chemicals

Oligonucleotides were synthesized on a PS 250 CRUACHEM DNA synthesizer using the cyanoethyl phosphoroamidate method, and purified by gel filtration. The sequence of the double-stranded oligonucleotide used to detect the DNA-binding activities of NF- κ B is shown with the binding site underlined: 5'-GATCAGTTGAGGGGACTTTCCAGGC-3'. DMEM, RPMI 1640, foetal bovine serum, penicillin/streptomycin and autocamide-3 were obtained from Gibco BRL (Grand Island, NY, U.S.A.). [α - ^{32}P]dATP (3000 Ci mmol^{-1}), [^{32}P]orthophosphate, horseradish peroxidase-conjugated anti-mouse and anti-rabbit antibodies, and the ECL detection agent were purchased from Amersham Pharmacia Biotech (Piscataway, NJ, U.S.A.). Ro31-8220 and KN-93

were purchased from Calbiochem (La Jolla, CA, U.S.A.). D609 was from Biomol (Plymouth Meeting, PA, U.S.A.). Rabbit polyclonal antibodies specific for p65 NF- κ B, p50 NF- κ B, I κ B α , I κ B β , IKK α , IKK β , ERK2, and p38 MAPK, and protein A/G beads were from Santa Cruz Biotechnology (Santa Cruz, CA, U.S.A.). Plasmid of pGEX-I κ B α (amino acids 5–55) was provided by Dr Frank S. Lee (Pennsylvania Medical Center, PA, U.S.A.). Plasmids of pGEX-p65 (354–551) and pGEX-p65 (point mutant of amino acid Ser⁵³⁶ substitution by Ala⁵³⁶) were a kind gift from Dr H. Sakurai (Tanabe Seiyaku, Osaka, Japan). All materials for SDS-PAGE were obtained from Bio-Rad Laboratories (Hercules, CA, U.S.A.). All other chemicals were obtained from Sigma (St. Louis, MO, U.S.A.).

Statistical analysis

Values are expressed as the mean \pm standard error of the mean (s.e.mean) of at least three experiments. Analysis of variance (ANOVA) was used to assess the statistical significance of the differences, and a *P* value less than 0.05 was considered statistically significant.

Results

Inhibition of LPS-induced NF- κ B activation by D609, Ro 31-8220, PD 098059, and SB 203580

Our previous results showed that LPS can induce NF- κ B activation as well as I κ B α phosphorylation and degradation, and in turn cause iNOS, COX-2, and IL-6 gene expression in murine J774 macrophages. Moreover, activation of NF- κ B is indispensable for these LPS actions (Chen *et al.*, 1998; Chen & Lin, 1999; 2000).

To investigate the regulatory roles of phosphatidylcholine-phospholipase C (PC-PLC), PKC, ERK, and p38 MAPK in LPS-induced NF- κ B activation, D609 (a selective PC-PLC inhibitor) (Muller-Decker, 1989), Ro 31-8220 (a selective PKC inhibitor) (Keller & Niggli, 1993), PD 098059 (a selective mitogen-activated protein kinase kinase (MEK) inhibitor) (Dudley *et al.*, 1995), and SB 203580 (a selective p38 MAPK inhibitor) (Lee *et al.*, 1994) were examined. When J774 macrophages were treated for 30 min with LPS, the nuclear translocation of NF- κ B was evidenced by *in vitro* binding to a specific DNA sequence. First, we performed antibody gel super-shift assays to determine whether the LPS-induced DNA-protein complex contained p50 (NF κ B1) and p65 (RelA) subunits. Upon addition to nuclear extracts prepared from LPS-treated cells, p50-specific and p65-specific antibodies caused super-shifts of the DNA-protein complexes, p50/p65 heterodimer, and p50/p50 homodimer (Figure 1, arrowheads), suggesting the existence of two types of NF- κ B complexes. When J774 cells were pretreated for 20 min with (μM): pyrrolidine dithiocarbamate (PDTC) 50, D609 30, Ro 31-8220 1, PD 098059 30, or SB 203580 3, LPS-induced NF- κ B activation was inhibited by $40 \pm 7\%$, $64 \pm 5\%$, $69 \pm 11\%$, $66 \pm 6\%$, and $72 \pm 5\%$ ($n = 3$), respectively (Figure 1). At the concentrations used, none of these inhibitors significantly affected the basal level of NF- κ B. These results suggest that PKC, ERK, and p38 MAPK are all involved in the upstream signal pathways which mediate LPS-elicited

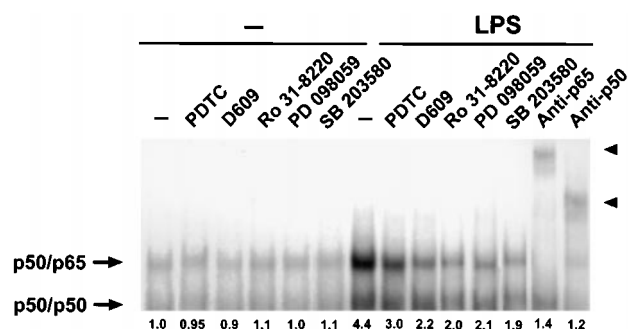


Figure 1 Effects of pharmacological agents on LPS-induced NF- κ B activation. J774 macrophages were preincubated with a vehicle (in μ M): PDTC 50, D609 30, Ro 31-8220 1, PD 098059 30, or SB 203580 3, for 20 min followed by stimulation with LPS ($1 \mu\text{g ml}^{-1}$) for another 30 min. Following incubation, nuclear extracts were prepared and subjected to EMSA as described in the Methods section. The NF- κ B species in the DNA-protein complex was determined by a super-shift with inclusion of $4 \mu\text{g}$ of the anti-p65 or anti-p50 antibody in the nuclear extract of LPS-stimulated cells as indicated by arrowheads. The results are representative of three independent experiments.

NF- κ B activation. Furthermore, this suggests that PC-PLC-derived diacylglycerol is a major component for LPS-induced PKC activation.

Effects of pharmacological inhibitors on $\text{I}\kappa\text{B}\alpha$ degradation

NF- κ B is generally retained in the cytoplasm of unstimulated cells by interaction with $\text{I}\kappa\text{B}\alpha$ and $\text{I}\kappa\text{B}\beta$. Upon response to most NF- κ B-inducing signals, both $\text{I}\kappa\text{B}\alpha$ and $\text{I}\kappa\text{B}\beta$ are targeted for IKK-specific phosphorylation, followed by proteasome-dependent degradation, resulting in NF- κ B dissociation from $\text{I}\kappa\text{B}\alpha$, as well as its nuclear translocation and DNA binding. To investigate the underlying mechanism for NF- κ B inhibition caused by kinase inhibitors, we next studied whether the PKC-, ERK-, and p38 MAPK-dependent pathways are essential for LPS-mediated $\text{I}\kappa\text{B}$ degradation. As shown in Figure 2a, treatment with LPS for 10 min resulted in $\text{I}\kappa\text{B}\alpha$ degradation, which reached the maximum extent at 30 min and began to recover after that because of re-synthesis of $\text{I}\kappa\text{B}\alpha$. LPS also induced $\text{I}\kappa\text{B}\beta$ degradation, albeit with a slower onset at 30 min and a delayed maximal effect at 60 min (Figure 2a). These results are consistent with previous findings showing that LPS exhibits a delayed kinetic on the degradation of $\text{I}\kappa\text{B}\beta$ compared to that on $\text{I}\kappa\text{B}\alpha$ (Velasco *et al.*, 1997). We next examined the effects of these inhibitors on LPS-induced $\text{I}\kappa\text{B}\alpha$ degradation. As shown in Figure 2b, PDTC (50 μM), D609 (30 μM), Ro 31-8220 (1 μM), and PD 098059 (30 μM), but not SB 203580 (3 μM), markedly reduced LPS induction of $\text{I}\kappa\text{B}\alpha$ degradation. At the concentrations used, none of these inhibitors affected the basal level of $\text{I}\kappa\text{B}\alpha$ protein. These results suggest the crucial roles played by PKC and ERK, but not p38 MAPK, in the upstream signal cascades required for LPS induction of $\text{I}\kappa\text{B}\alpha$ degradation.

Time-dependent IKK activation by LPS and UTP

Our previous study demonstrated that P2Y_6 receptor stimulation by UTP triggers a Ca^{2+} -dependent pathway

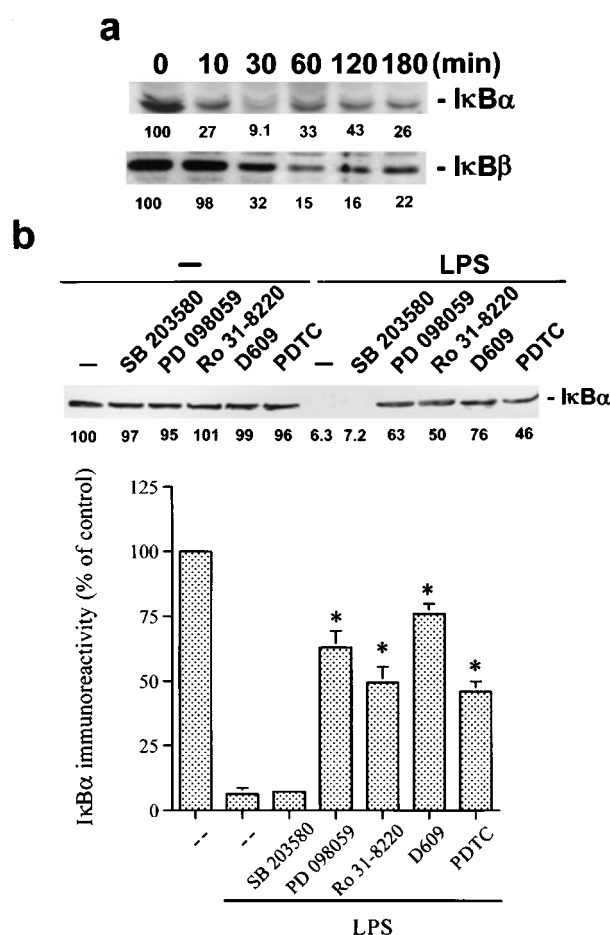


Figure 2 Effects of protein kinase inhibitors on LPS-induced $\text{I}\kappa\text{B}$ degradation. (a) Time course analysis of $\text{I}\kappa\text{B}\alpha$ and $\text{I}\kappa\text{B}\beta$ degradation induced by LPS. Following incubation for different periods with LPS ($1 \mu\text{g ml}^{-1}$), $\text{I}\kappa\text{B}\alpha$ and $\text{I}\kappa\text{B}\beta$ degradations were determined by immunoblotting with $\text{I}\kappa\text{B}\alpha$ - or $\text{I}\kappa\text{B}\beta$ -specific antibody. (b) Cells were preincubated with vehicle (in μM): PDTC 50, D609 30, Ro 31-8220 1, PD 098059 30, or SB 203580 3, for 20 min and then incubated with LPS ($1 \mu\text{g ml}^{-1}$) for an additional 30 min. Measurement of $\text{I}\kappa\text{B}\alpha$ degradation was as described above. The traces are representative of three experiments with similar results, which are shown as the mean \pm s.e.mean. * $P < 0.05$ as compared to the control LPS response without inhibitor pretreatment.

that synergizes LPS-induced NF- κ B activation by increasing phosphorylation and degradation of $\text{I}\kappa\text{B}\alpha$ (Chen *et al.*, 1998). To determine whether this enhanced $\text{I}\kappa\text{B}$ phosphorylation by UTP results from upstream IKK activation, we directly measured IKK complex kinase activity by immunoprecipitating $\text{IKK}\alpha$ and $\text{IKK}\beta$ from J774 cells that were either activated or not by UTP and/or LPS. As shown in Figure 3, treatment of J774 cells with LPS ($1 \mu\text{g ml}^{-1}$) elicited a marked (about a 15 fold activity increase with 20 min of incubation) and sustained (lasting for at least 3 h) activation of the IKK complex (Figure 3a). Compared to the LPS response, UTP (100 μM) elicited a slight and transient IKK activation (Figure 3a).

To explore the mechanism responsible for UTP-induced IKK activation, two pharmacological agents, thapsigargin and phorbol 12-myristate 13-acetate (PMA), which respectively mimic PI-PLC-transduced Ca^{2+} - and PKC-dependent

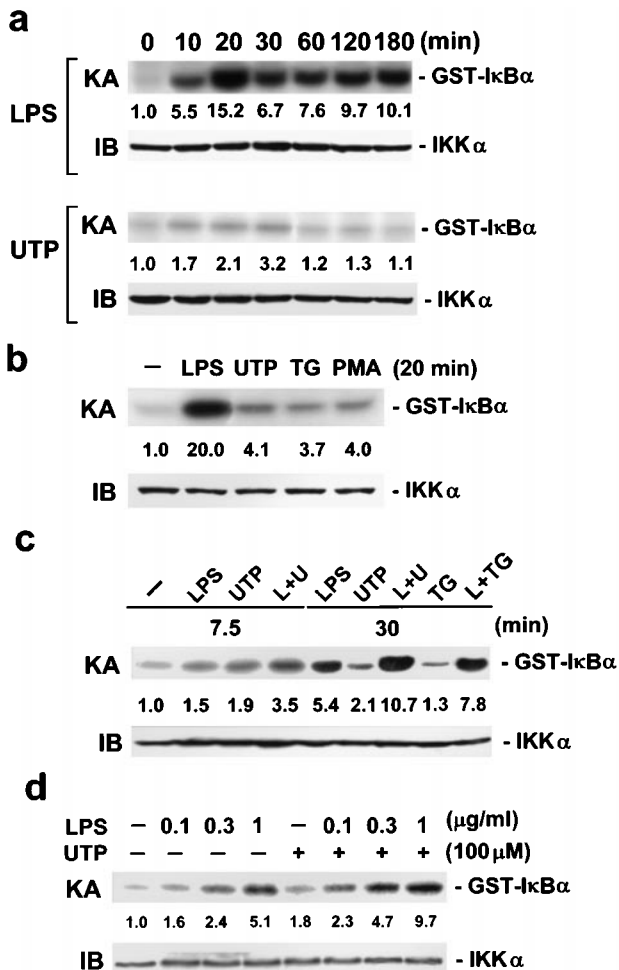


Figure 3 IKK activation by LPS and UTP. Cells were, as indicated, treated with $1 \mu\text{g ml}^{-1}$ LPS, $100 \mu\text{M}$ UTP, 30 nM thapsigargin (TG), $1 \mu\text{M}$ PMA (a–c) or 0.1 – $1 \mu\text{g ml}^{-1}$ LPS (d) for different intervals, and the cell lysates were then immunoprecipitated with antibodies specific for IKKs. One set of immunoprecipitates was subjected to kinase assay (KA) as described in the Methods section using GST-IκBα (5–55) as a substrate (a, b, c and d, top). The multiples of induction of IKK activity under each stimulation condition are shown. The other set of immunoprecipitates was subjected to SDS-PAGE and analysed by immunoblotting (IB) with anti-IKKα antibody (a, b, c and d, bottom). Equal amounts of the immunoprecipitated kinase complex present in each kinase assay were confirmed by immunoblotting for IKKα. Shown are representative results from three independent experiments.

signalling cascades, were investigated. Treatment with thapsigargin (30 nM) or PMA ($1 \mu\text{M}$) for 20 min moderately increased IKK activity to a similar extent as did UTP (Figure 3b), suggesting that both bifurcating signal pathways of PI-PLC, Ca^{2+} and PKC, participate in UTP activation of IKK. The combined effects of UTP and LPS on IKK activity, as shown in Figure 3c, also indicate their synergistic action, with a more marked response seen at 30 min than at 7.5 min . Consistent with UTP effect, thapsigargin potentiation of LPS-induced IKK activity was seen (Figure 3c). The concentration-dependent IKK activation by LPS and potentiation by UTP was shown in Figure 3d. UTP at $100 \mu\text{M}$ elicited dramatic potentiation of LPS response at 0.3 and $1 \mu\text{g ml}^{-1}$.

Roles of PKC, ERK, and CaMK in LPS- and UTP-induced IKK activation

To support our suggestions that PKC and ERK are involved in LPS induction of IκB degradation and that CaMK participates in UTP potentiation of IκB degradation, we examined the effects of pharmacological inhibitors on IKK activation. As shown in Figure 4a, PDTC ($50 \mu\text{M}$), Ro 31-8220 ($1 \mu\text{M}$), and PD 098059 ($30 \mu\text{M}$) respectively reduced LPS-stimulated IKK activities by $65 \pm 6\%$, $77 \pm 4\%$, and $42 \pm 5\%$ ($n=3$), and UTP-stimulated IKK activities by $70 \pm 4\%$, $68 \pm 3\%$, and $56 \pm 2\%$ ($n=3$). Consistent with its inhibition of LPS-induced NF-κB activation and IκB degradation, D609 ($30 \mu\text{M}$) decreased LPS-induced IKK activity by $54 \pm 7\%$ ($n=3$), while it did not change IKK activation by UTP. On the contrary, upon cell treatment with $10 \mu\text{M}$ KN-93 (a selective CaMK inhibitor) (Sumi *et al.*, 1991), UTP-induced, but not LPS-induced, IKK activity was inhibited by $51 \pm 5\%$ ($n=3$) (Figure 4a). BAPTA/AM ($30 \mu\text{M}$), an intracellular calcium chelator, completely inhibited UTP-induced IKK activity, but did not affect LPS-induced IKK activity. SB 203580 ($3 \mu\text{M}$), on the other hand, did not affect IKK activation induced by either stimulus. Consistent with the individual attenuation of inhibitors, the synergistic IKK stimulation by LPS and UTP was sensitive to inhibition by Ro 31-8220, KN-93, PD 098059, Go 6976 (an inhibitor of classical PKC) and BAPTA/AM (Figure 4b). These results suggest that PKC

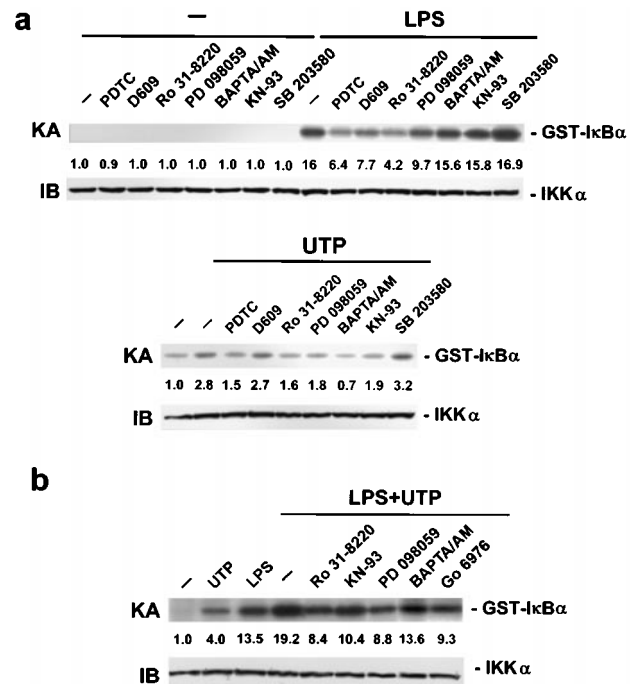


Figure 4 Effects of protein kinase inhibitors on LPS- and UTP-induced IKK activation. Cells were preincubated with vehicle ($\text{in } \mu\text{M}$): PDTC 50 , D609 30 , Ro 31-8220 1 , PD 098059 30 , BAPTA/AM 30 , KN-93 10 , SB 203580 3 or Go 6976 1 , for 20 min before the addition of LPS ($1 \mu\text{g ml}^{-1}$), UTP ($100 \mu\text{M}$), or both for another 20 min . Cell lysates were then immunoprecipitated with IKKs antibodies. Both kinase assays and IKKα immunoblotting were analysed as described in Figure 3. The multiples of induction of IKK activity under each stimulation condition are shown. The typical traces shown are representative of three experiments.

and ERK converge the signal cascades triggered by LPS and UTP to activate IKK, and that the Ca^{2+} -dependent CaMK pathway also participates in the UTP response.

LPS and UTP activation of ERK and p38 MAPK

To directly confirm the crucial roles of ERK and p38 MAPK in NF- κ B activation, we determined both kinase activities in response to LPS and UTP stimulation. The results using MBP as a kinase substrate indicated that LPS ($1 \mu\text{g ml}^{-1}$) and UTP ($100 \mu\text{M}$) stimulated ERK and p38 MAPK in a time-dependent manner, which occurred as early as 5–10 min of incubation and reached their peak responses at 30 min (Figure 5a,b). The ERK and p38 MAPK activity caused by LPS and UTP was respectively attenuated by the

presence of $30 \mu\text{M}$ PD 098059 (Figure 5c) and $3 \mu\text{M}$ SB 203580 (Figure 5d).

KN-93 inhibits UTP-induced CaMK activity

To confirm CaMK playing a major role in IKK potentiation by UTP, we determined CaMK activity following UTP treatment. The results indicated that the autonomous activity of CaMK was rapidly increased by $100 \mu\text{M}$ UTP treatment for 10 s (from $16.9 \pm 2.7\%$ of total CaMK activity in control cells to $25.8 \pm 0.5\%$ in UTP-treated cells). The CaMK activity stimulated by UTP was abrogated by the presence of $10 \mu\text{M}$ KN-93 (Figure 6).

LPS and UTP induction of p65 phosphorylation

In addition to $\text{I}\kappa\text{B}\alpha/\beta$ degradation which causes nuclear translocation of NF- κ B, post-translational modification of p65 NF- κ B subunit, such as phosphorylation, represents another mechanism for controlling NF- κ B activity. A recent study has shown that the endogenous IKK complex can phosphorylate $\text{I}\kappa\text{B}\alpha$ as well as p65 NF- κ B subunit in TNF- α -stimulated HeLa cells (Sakurai *et al.*, 1999). The serine residue at amino acid 536 was demonstrated to be the phosphorylation target of IKK (Sakurai *et al.*, 1999). To establish whether the UTP increase of LPS-induced IKK activation contributes to p65 phosphorylation, we performed two experiments. First, direct detection of p65 phosphorylation by *in vivo* [^{32}P] metabolic labeling plus immunoprecipitation analysis showed that both LPS ($1 \mu\text{g ml}^{-1}$) and UTP ($100 \mu\text{M}$) elicited p65 phosphorylation within 30 min and that both responded in an additive manner at 30 min (Figure 7a). Second, we found that the above results showing the synergistic IKK stimulation by LPS and UTP (Figures 3c and 4b) could also be reflected in p65 phosphorylation. Using GST-p65 (354–551) recombinant protein as the *in vitro* kinase substrate, LPS plus UTP showed synergistic effects on IKK activities as compared to the extent caused by each individually. Moreover, the recombinant protein with the point substitution of Ser⁵³⁶ with Ala was determined not to be

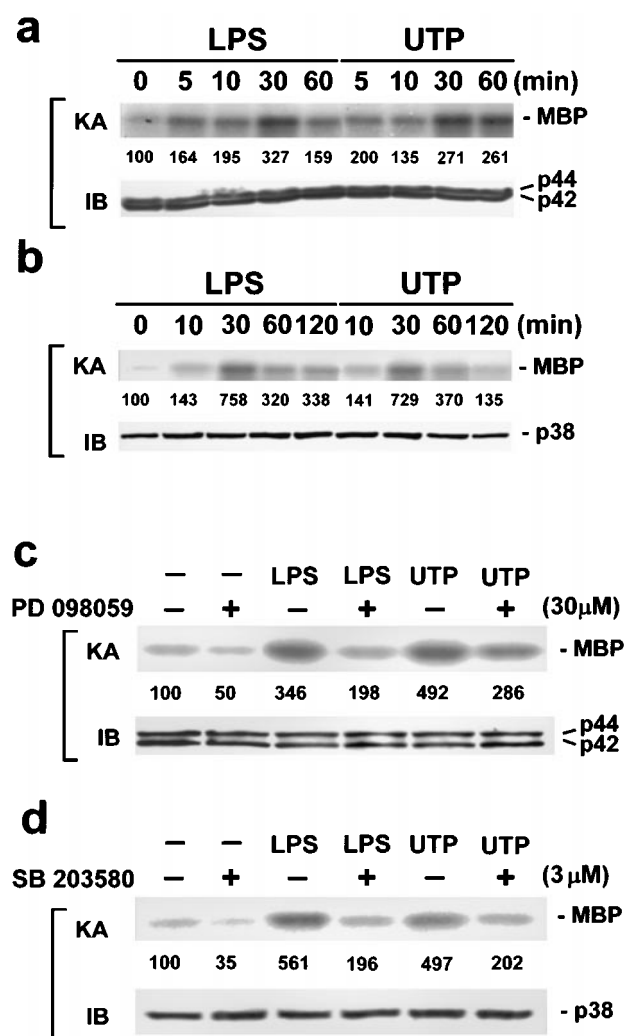


Figure 5 LPS- and UTP-induced ERK and p38 MAPK activation. After cells were treated with $1 \mu\text{g ml}^{-1}$ LPS or $100 \mu\text{M}$ UTP for different intervals, cell lysates were immunoprecipitated with an antibody specific for ERK (a) or p38 MAPK (b). Both kinase assays and MAPK immunoblotting were analysed as described. In some experiments, cells were preincubated with vehicle, $30 \mu\text{M}$ PD 098059 (c) or $3 \mu\text{M}$ SB 203580 (d) for 20 min before the addition of LPS ($1 \mu\text{g ml}^{-1}$) or UTP ($100 \mu\text{M}$) for another 30 min. Both kinase assays and MAPK immunoblotting were analysed as described. Shown are representative results from three to four independent experiments.

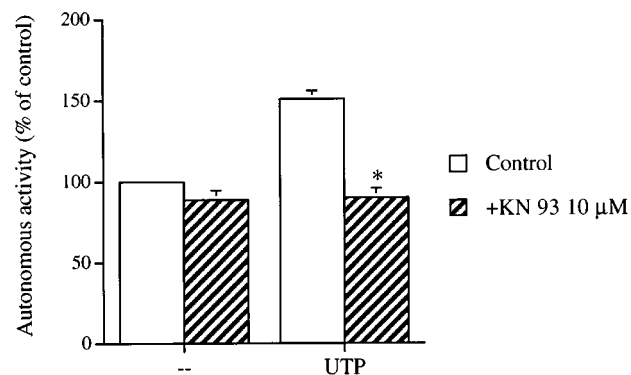


Figure 6 KN-93 inhibits UTP-induced CaMK activity in J774 cells. Cells were preincubated with vehicle, or $10 \mu\text{M}$ KN-93 for 20 min before the addition of UTP ($100 \mu\text{M}$) for another 10 s. The CaMK activity was assayed as described in the Methods section. Basal levels of autonomous activity was $44.1 \pm 1.4 \text{ pmol min}^{-1} \text{ mg}^{-1}$ ($n=3$). The data represent the mean \pm s.e. mean of three experiments performed in duplicate.

the IKK target, confirming the IKK phosphorylation site at Ser⁵³⁶ (Figure 7b).

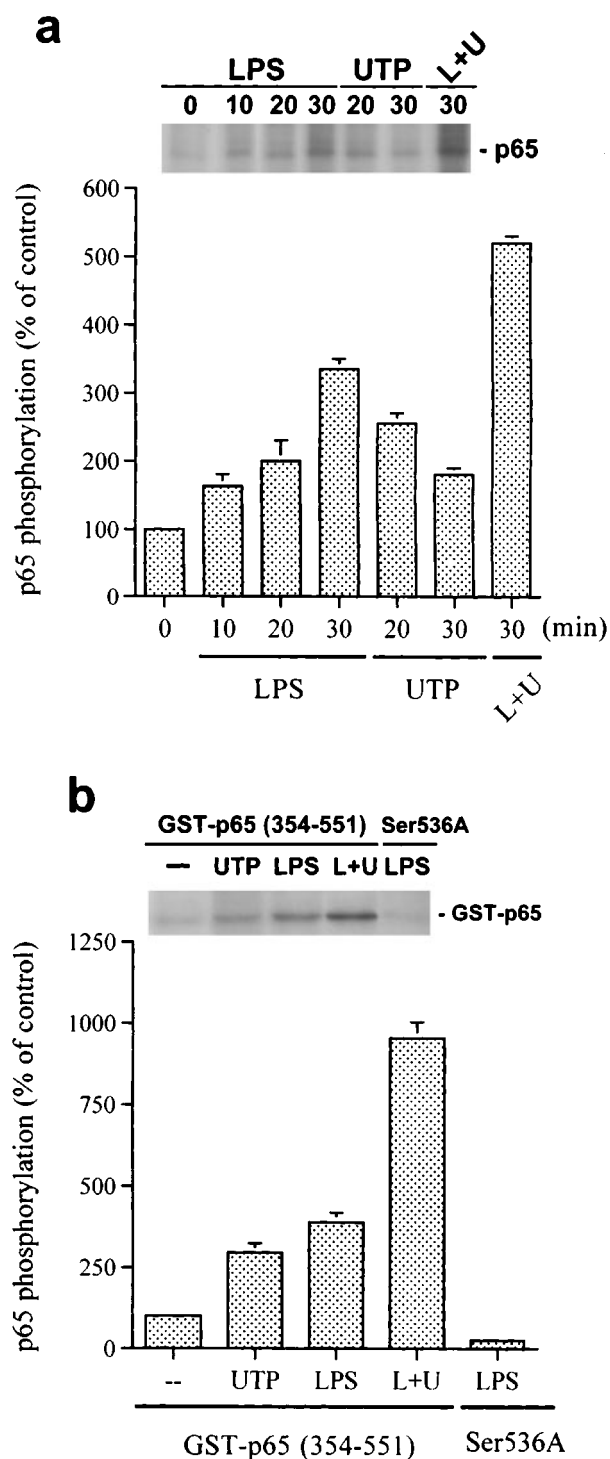


Figure 7 LPS and UTP mediation of p65 phosphorylation. (a) Cells loaded with [³²P]-orthophosphate were treated with LPS (1 $\mu\text{g ml}^{-1}$), UTP (100 μM), or both for the indicated time periods, and p65 was then immunoprecipitated and analysed by SDS-PAGE and autoradiography. (b) Cells were treated with LPS with or without UTP for 30 min, and cell lysates were then immunoprecipitated with antibodies specific for IKKs. The enzyme activity of IKK immunoprecipitates was assayed by using GST-p65 (354–551) or GST-p65 (Ser⁵³⁶ to Ala⁵³⁶) as a substrate. Traces are representative results from three independent experiments, which are shown as mean \pm s.e.mean.

LPS and TG induction of IKK activity in peritoneal macrophages

We next confirm that the calcium-dependent IKK potentiation observed in the J774 cell line is not cell type-dependent. Thapsigargin was previously shown to enhance LPS-elicited NO, prostaglandin E₂ (PGE₂), IL-6, and TNF- α release from peritoneal macrophages via a CaMK-dependent mechanism (Chen *et al.*, 2001). Using recombinant I κ B α as a substrate, we found that LPS (1 $\mu\text{g ml}^{-1}$) and thapsigargin (30 nM) increased peritoneal macrophage IKK activity with similar kinetics. Each stimulus elicited its maximal IKK activity at 20 min (Figure 8a). Although the thapsigargin response was less than that of LPS, thapsigargin also exerted a synergistic effect with LPS. This synergistic IKK activation was further evidenced by using GST-p65 (354–551) as a kinase substrate (Figure 8b). Likewise, the IKK activity caused by LPS plus thapsigargin was respectively attenuated by the presence of BAPTA/AM (30 μM), KN-93 (10 μM), Ro 31-8220 (1 μM), or PD 098059 (30 μM) (Figure 8c).

Discussion

NF- κ B is a ubiquitous transcription factor and plays a crucial role in regulating many genes in a variety of cell types. Although NF- κ B activation has been widely studied in many kinds of cytokine signalling, only a little is known about its activation by G protein-coupled receptors. Histamine (Hu *et al.*, 1999), platelet-activating factor (Kravchenko *et al.*, 1995), lysophosphatidic acid (Shahrestanifar *et al.*, 1999), thrombin (Mari *et al.*, 1994; Rahman *et al.*, 1999), bradykinin (Pan *et al.*, 1996; Xie, *et al.*, 2000), leukotriene B₄, fMet-Leu-Phe (Browning *et al.*, 1997), substance P (Lieb *et al.*, 1997), endothelin-1 (Gallois *et al.*, 1998), and carbachol (Li *et al.*, 1996) were reported to induce NF- κ B activation, while the upstream signalling molecules required for NF- κ B activation by these G protein-coupled receptors have not been clearly explored.

In J774 macrophages, UTP is known to bind the P2Y₆ receptors to stimulate PI breakdown, increase [Ca²⁺]_i, and activate PKC isoforms (Lin & Chen, 1997; Chen & Lin, 2000). In addition, UTP elicits the potentiation of LPS-induced iNOS (Chen *et al.*, 1998), COX-2 (Chen & Lin, 2000), IL-6 (Chen & Lin, 1999), and TNF- α (Chen *et al.*, 2001) gene expression. Notably our previous studies further indicated that CaMK-dependent NF- κ B potentiation primarily accounts for these potentiation effects of UTP (Chen *et al.*, 1998; 2001; Chen & Lin, 1999; 2000). To better understand the upstream mechanism for NF- κ B activation, in this study, we primarily focused on the biochemical assay of IKK, which is well known as a key element in the signal transduction cascade that leads to phosphorylation and degradation of I κ B α / β , and thereby subsequently induces NF- κ B nuclear translocation and transactivation. We have examined not only the effect of UTP on IKK activity but also the possible involvement of the PKC, ERK, p38 MAPK, and CaMK pathways in this respect. The present study shows the ability of the Gq-coupled receptor to activate the IKK complex, and the involvement of CaMK, PKC, and ERK in this action.

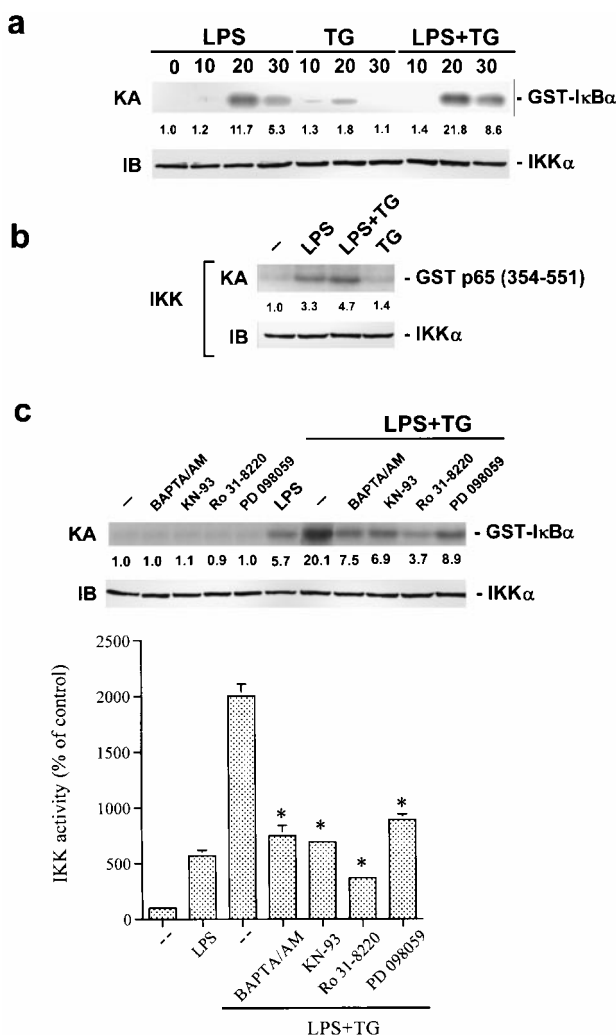


Figure 8 IKK activation by LPS and thapsigargin in peritoneal macrophages. (a) Cells were treated with $1 \mu\text{g ml}^{-1}$ LPS, 30 nM thapsigargin (TG), or both for different intervals, then cell lysates were immunoprecipitated with IKKs antibodies. Both kinase assay, as an index to GST-IκB α phosphorylation, and IKK immunoblotting were used to analyse IKK immunoprecipitates. (b) Cells were treated with LPS with or without thapsigargin for 20 min, then cell lysates were immunoprecipitated with IKK antibodies and assayed for kinase activity toward GST-p65 (354–551) phosphorylation. (c) Cells were preincubated with (μM): BAPTA/AM 30, KN-93 10, Ro 31-8220 1, or PD 098059 30, for 20 min before the addition of LPS with or without thapsigargin for another 20 min. Cell lysates were then immunoprecipitated as described. The typical traces are representative of three experiments, which are shown as mean \pm s.e.mean. * $P < 0.05$ as compared to the control response of LPS plus thapsigargin without inhibitor pretreatment.

Several studies have demonstrated the indispensability of the generation of reactive oxygen intermediates for IKK-dependent NF- κ B activation. Thus PDTC, a scavenger of reactive oxygen intermediates, has been widely used as an inhibitor of NF- κ B. In our experimental system, we show that PDTC is able to inhibit LPS-induced IKK activity, IκB degradation, and NF- κ B activation as well as UTP-induced IKK activity. All these results further confirm the involvement of reactive oxygen intermediates in IKK activation.

Our data provide evidence to show that the two downstream signal pathways of PI turnover, Ca^{2+} increase and PKC activation, mediate UTP activation of IKK. BAPTA/AM (an intracellular Ca^{2+} chelator), KN-93 (a selective CaMK inhibitor), Ro 31-8220 and Go 6976 (two selective PKC inhibitors) all effectively reduce the UTP response. Our previous study has shown that the increased intracellular Ca^{2+} concentration by UTP in macrophages is neutralized by BAPTA/AM (Chen *et al.*, 2001). Extending this finding, we showed the ability of UTP to induce a KN-93 sensitive CaMK activity in this study. The results that IKK activation by thapsigargin in peritoneal macrophages is also attenuated by BAPTA/AM, KN-93 and Ro 31-8220 further support the key roles of Ca/CaMK and PKC in mediating IKK activation. Although thapsigargin is not a stimulator of PI turnover, it can somewhat mimic downstream signals of PI turnover *via* its inhibition of endoplasmic reticulum Ca^{2+} -ATPase, its increase in $[\text{Ca}^{2+}]_i$, and its activation of conventional PKC isoforms (Thastrup, 1990). The involvement of PKC is supported by the moderate IKK activation caused by the potent PKC activator, PMA, and the inhibitory abilities of the two selective PKC inhibitors, Ro 31-8220 and Go 6976, in reducing IKK activation. Although IKK activity is increased by UTP as shown in this study, UTP in the same cell system cannot cause any significant changes in IκB degradation or NF- κ B activation, which has previously been reported (Chen *et al.*, 1998). We think this inconsistency might be explained by the insufficiency of the weak and transient IKK activity to trigger IκB phosphorylation.

NF- κ B activation is one of the most important and phylogenetically conserved cellular signalling transduction pathways implicated in the innate immune response and pathology of microbial pathogen LPS (Hoffmann *et al.*, 1999). LPS-induced IKK activation pathways have been intensively studied, and recent genetic and biochemical experiments have highlighted the critical roles of several signal mediators in LPS signalling and have directed their sequential signalling towards NF- κ B activation as the CD14/TLR \rightarrow MyD88 \rightarrow IRAK \rightarrow TRAF6 \rightarrow TAK1 \rightarrow NIK \rightarrow IKK complex (Kirschning *et al.*, 1998). However, besides these well-characterized signal pathways, LPS is suggested to act through different signal pathways on the same integrator IKK. This suggestion is primarily based on the finding that LPS is still able to elicit NF- κ B activity in MyD88 $^{-/-}$ macrophages (Kawai *et al.*, 1999).

Several pieces of evidence obtained from recent reports have shown the role of PKC in IKK regulation. GF 109203X, an inhibitor of conventional and novel PKC isoforms, abrogates PMA and ionomycin activation of the IKK complex in Jurkat and primary CD3 $^{+}$ T lymphocytes (Trushin *et al.*, 1999). Transient overexpression studies have shown that PKC ϵ (Genot *et al.*, 1995) and PKC α (Trushin *et al.*, 1999) induce, while the dominant negative mutant of PKC ζ (Lozano *et al.*, 1994) impairs, NF- κ B-dependent reporter gene activation. In J774 macrophages, we have shown that PKC activation is required for LPS induction of iNOS, COX-2, and IL-6 gene expressions (Chen *et al.*, 1998; Chen & Lin, 1999; 2000). In the present study, the selective PKC inhibitor, Ro 31-8220, inhibits LPS-induced NF- κ B activation, IκB degradation, and IKK activity, strongly

indicating that PKC is a supplementary signalling pathway for LPS-mediated NF- κ B activation.

PKC is usually activated by the physiological activator, diacylglycerol, which can be generated by the enzymatic action of PI-PLC and PC-PLC. Recent studies have demonstrated that LPS-mediated diacylglycerol formation and PKC activation result from PC-PLC pathway and is inhibited by D609 (a PC-PLC inhibitor) (Tschaikowsky *et al.*, 1994; Monick *et al.*, 1999). In this study we found that D609 is able to inhibit LPS-induced NF- κ B activation, I κ B degradation, and IKK activation, but has no effects on UTP. This result supports our previous report indicating that LPS-induced NO, PGE₂, and IL-6 releases are sensitive to treatment with D609 but not with U73122 (a PI-PLC inhibitor). On the contrary, UTP potentiation of LPS action is resistant to D609 but sensitive to U73122 (Chen *et al.*, 1998). Thus, although coupling to different PLC systems and catalysing different phospholipid substrates, both LPS and UTP generate diacylglycerol and activate PKC. The fact that LPS is unable to activate PI turnover or increase [Ca²⁺]_i (data not shown) together with the finding that its signalling (Figure 4a) and functional responses (Chen *et al.*, 1998; Chen & Lin, 2000) are unaffected by BAPTA/AM and KN-93 all confirm its activation of PC-PLC but not PI-PLC.

Pharmacological approaches have demonstrated the requirement of p38 MAPK and/or ERK for NF- κ B transactivation in response to LPS (Chen *et al.*, 1999), TNF- α (Bergmann *et al.*, 1998; Vanden Berghe *et al.*, 1998), and IL-1 (Bergmann *et al.*, 1998). Although both MAPKs have been suggested to act as necessary and cooperative machinery for NF- κ B transactivation, the mechanism is not clear yet. In the present study, we found that PD 098059 (a MEK inhibitor) attenuates not only LPS-mediated IKK activation, I κ B degradation, and NF- κ B activation, but also UTP-induced

IKK activation. These results suggest that ERK also plays a role in IKK activation in response to LPS and UTP. In this study, although the p38 MAPK inhibitor, SB 203580, inhibited NF- κ B activation, it had no effects on IKK-dependent I κ B degradation. Thus, as previously suggested (Bergmann *et al.*, 1998), an I κ B degradation-independent mechanism may contribute to p38 MAPK-linked NF- κ B activation.

In addition to the NIK/IKK/I κ B cascade, recent data have shown that different NF- κ B-inducing stimuli such as TNF- α and IL-1 appear to cause p65 RelA phosphorylation at unique sites prior to its nuclear translocation, and to positively regulate NF- κ B-dependent transcription (Vanden Berghe *et al.*, 1998; Sakurai *et al.*, 1999; Jefferies & O'Neill, 2000). Protein kinases such as IKK (Sakurai *et al.*, 1999), PKA (Zhong *et al.*, 1998), p38 MAPK (Vanden Berghe *et al.*, 1998; Jefferies & O'Neill, 2000), ERK (Vanden Berghe *et al.*, 1998; Jefferies & O'Neill, 2000), and PKC (Lozano *et al.*, 1994) have all been shown to phosphorylate p65. In this study, we show that IKK activation by LPS and UTP can target p65 phosphorylation at Ser⁵³⁶, and p65 phosphorylation induced by both stimuli *in vivo* might contribute to their coordinated increase in κ B gene transcription.

Taken together, this report demonstrates that Gq protein-coupled receptor activation can activate IKK and potentiate LPS-stimulated NF- κ B activation. Not only do PKC and ERK play regulatory roles in LPS activation of IKK/I κ Bs/NF- κ B signalling, but the increase in intracellular calcium and subsequent CaMK activation also play a major role in UTP potentiation of the LPS response.

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