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Redox/ROS regulation of lipopolysaccharide-induced mitogen-activated protein kinase (MAPK) activation and MAPK-mediated TNF- α biosynthesis

*^{,1,2}J.J. Haddad & ¹S.C. Land

¹Oxygen Signalling Group, Centre for Research into Human Development, Tayside Institute of Child Health, Faculty of Medicine, Ninewells Hospital and Medical School, University of Dundee, DD1 9SY

1 Redox and ROS regulation of MAPK-mediated TNF- α biosynthesis is not well characterized. It was hypothesized that the involvement of the MAPK pathway in regulating LPS-mediated TNF- α secretion is redox-dependent, NF- κ B-sensitive and attenuated by *N*-acetyl-L-cysteine (NAC) and other antioxidants.

2 In alveolar epithelial cells, LPS induced a time- and dose-dependent phosphorylation of MAPK^{p38}. This was associated with the activation of MAPK-activated protein kinase, which phosphorylated the small heat-shock protein, Hsp27.

3 MAPK^{p38} inhibition (SB-203580) abrogated LPS-induced TNF- α production. MAPK^{ERK} blockade (PD-98059) attenuated TNF- α secretion, an effect synergistically amplified in the presence of SB-203580.

4 Regulation of NF- κ B by selective inhibitors revealed that this pathway is partially involved in regulating LPS-mediated TNF- α secretion. Whereas the proteasome inhibitor, MG-132, had no effect on LPS-mediated TNF- α production, CAPE, sulfasalazine and SN-50, a cell-permeant NF- κ B inhibitor, attenuated but did not abrogate TNF- α biosynthesis.

5 LPS up-regulated ROS, an effect abrogated by 4'-hydroxy-3'-methoxy-acetophenone and NAC, which reduced TNF- α secretion, induced the accumulation of GSH, reduced the concentration of GSSG, and blockaded the phosphorylation/activation of MAPK^{p38} pathway.

6 ROS induced MAPK^{p38} phosphorylation and selective antioxidants, including the permeant GSH precursor, γ -GCE, reduced ROS-dependent MAPK^{p38} phosphorylation.

7 These results indicate that the MAPK pathway and MAPK-mediated regulation of $TNF-\alpha$ production is redox-dependent, GSH-mediated and requires, at least in part, a NF- κ B/ROS-sensitive mechanism.

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Abbreviations: AEBSF, 4-(2-aminoethyl)]-benzene sulphonyl fluoride-HCl; ANOVA, analysis of variance; ARE, AU-rich element; BSA, bovine serum albumin; DMEM, Dulbecco's modified Eagle medium; DMSO, dimethyl sulphoxide; DTT, dithiothreitol; ELISA, enzyme-linked immuno-sorbent assay; ERK, extracellular signal-regulated kinase; fATII, foetal alveolar type II epithelial cells; FCS, foetal calf serum; Glutathione (GSH), L-γ-glutamyl-L-cysteinyl-glycine; glutathione (GSSG), oxidized glutathione disulphide; H₂O₂, hydrogen peroxide; HBSS, Hanks' balanced salt solution; Hsp27, heat-shock protein 27; HRP, horseradish peroxidase; IκB, inhibitory-κB; IKK, IκB kinase; IL, interleukin; LPS, lipopolysaccharide; LBP, lipopolysaccharide binding protein; MKK, MAPK kinase; MAPK, mitogen-activated protein kinase; O₂⁻⁻, superoxide anion; 'OH, hydroxyl radical; PBS, phosphate buffered saline; PGE₂, prostaglandin-E₂; redox, reduction-oxidation; RK, reactivating kinase; ROS, reactive oxygen species; SAPK, stress-activated protein kinase; SDS – PAGE, sodium-dodecyl polyacrylamide gel electrophoresis; Thr, threonine; TMB, 3,3,5,5-tetramethyl-benzidine dihydrochloride; TNF- α , tumour necrosis factor- α ; TRX, thioredoxin; Tyr, tyrosine

Introduction

Many extracellular stimuli, including pro-inflammatory cytokines and other inflammatory mediators (Herlaar & Brown, 1999), elicit specific cellular responses through the activation of mitogen-activated protein kinase (MAPK) signalling pathways (Marshall, 1995; Garrington & Johnson, 1999). MAPKs are proline-targeted serine-threonine kinases that transduce environmental stimuli to the nucleus and they themselves are activated by upstream MAPK kinases (MAPKKs) on both threonine and tyrosine residues within an 'activation loop' (Su & Karin, 1996; Garrington & Johnson, 1999). Once activated, MAPKs can phosphorylate and activate other kinases or nuclear proteins, including potential transcription factors, such as NF- κ B, and substrates (Stein *et al.*, 1997). The novel

^{*}Author for correspondence; E-mail: haddadj@anesthesia.ucsf.edu. ²Current address: Neuroscience Research Laboratory, Department of Anesthesia & Perioperative Care, University of California, Medical Centre S261, San Francisco, California, CA 94143, U.S.A.

mammalian reactivating protein kinase (p38/RK) MAPKs and ERK are stress-activated protein kinases (SAPK) that mediate responses to cellular stresses such as u.v. irradiation, osmotic imbalance, heat shock, DNA damage, bacterial products such as lipopolysaccharide (LPS), and inflammatory signals (Lee *et al.*, 1994; Wang *et al.*, 1997). Furthermore, inflammatory mediators, such as cytokines, including tumour necrosis factor (TNF)- α , activate the MAPK pathway in several cell types (Raingeaud *et al.*, 1995; Moriguchi *et al.*, 1997; Zu *et al.*, 1998). Of note, MAP kinase has been recently implicated in regulating pro-inflammatory cytokine biosynthesis (Ballard-Croft *et al.*, 2001; Lee *et al.*, 1994) and transcription (Rutault *et al.*, 2001).

The tripeptide L-y-glutamyl-L-cysteinyl-glycine, or glutathione (GSH), a ubiquitous thiol, plays a major role in maintaining intracellular reduction-oxidation (redox) equilibrium in the lung (Meister, 1988; Hayes & McLellan, 1999; Rahman, 1999; 2000; Haddad & Land, 2000a; Haddad et al., 2000). The cysteinyl moiety of GSH provides the reactive thiol as a functional element responsible for the diverse properties of glutathione, including (i) an antioxidant potential mediated by the peroxidase coupled reaction, (ii) regulation of cellular sulphydryl status and redox equilibrium, (iii) governing pathways in neuro-immune-endocrine interactions as an immunopharmacological thiol, and (iv) regulation of the expression/activation of redox-sensitive transcription factors induced by stress-evoked responses (Dröge et al., 1994; Hayes & McLellan, 1999; Haddad et al., 2000). The pivotal role of redox cycle in maintaining the integrity of the biological system in the face of oxidative stress and other challenges is, therefore, of particular relevance. The immunopharmacological potential assigned to glutathione (Thompson et al., 1985; Dröge et al., 1994), furthermore, stems from established observations. Interleukin (IL)-1 induced responses in mesangial cells, for instance, occurred through modulating redox equilibrium (Rovin et al., 1997). In addition, reactive oxygen species (ROS) signalling pathways regulating the transcription of IL-4 (Jeannin et al., 1995), IL-6, IL-8 (Gosset et al., 1999) and TNF-a (Neuschwander-Tetri et al., 1996; Gosset et al., 1999) were mediated through a thiol-dependent mechanism. Interestingly, antioxidants (Matsumoto et al., 1998; Reimund et al., 1998; Barrett et al., 1999; Haddad et al., 2001a) and glutathione precursors (Jeannin et al., 1995; Gosset et al., 1999; Pena et al., 1999; Haddad et al., 2001c) have been shown to down-regulate cytokine synthesis, activation and downstream processes. In this respect, Nacetyl-L-cysteine (NAC), an antioxidant and a GSH precursor (Bernard, 1991; Haddad et al., 2000), has been shown to ameliorate cytokine transcription and synthesis (Matsumoto et al., 1998; Tsuji et al., 1999; Hashimoto et al., 2001), and suppress ROS-mediated lung injury (Bernard, 1991).

Redox regulation of MAPK^{p38}- and MAPK^{ERK}-mediated cytokine transcription and biosynthesis has not been well characterized. It has been shown previously that thioredoxin (TRX), a redox control protein, negatively regulated MAPK^{p38} kinase activation and MAPK^{p38}-mediated IL-6 expression (Hashimoto *et al.*, 1999a, b). In addition, Hashimoto *et al.* (2001) recently reported that NAC attenuated TNF- α -induced MAPK^{p38} activation and the downstream IL-8 dependent pathway. One mechanism involved in the generation of ROS in a variety of cells is the oxidative burst, a process catalysed by the multicomplex

enzyme NADPH oxidase; MAPK^{p38} has been implicated in regulating this complex and subsequently coupled intracellular redox signalling with the oxidative burst-dependent regulation of pro-inflammatory signals (Ridley *et al.*, 1997; Nick *et al.*, 1999; Partrick *et al.*, 2000). However, redox regulation of LPS-induced MAPK-mediated TNF- α biosynthesis in the alveolar epithelium has yet to be ascertained and the underlying signalling mechanism was unravelled. Therefore, the aim of the present study was to investigate the role that glutathione and antioxidants play in regulating MAPK phosphorylation/activation, and to determine whether ROS and redox equilibrium are involved in MAPK-dependent regulation of TNF- α biosynthesis.

Methods

Chemicals and reagents

Unless otherwise indicated, chemicals of the highest analytical grade were purchased from Sigma-Aldrich (Dorset, England, U.K.). The pyridinyl imidazole SB-203580, a specific inhibitor of MAPK^{p38} (Lee et al., 1994) and PD-98059, a specific inhibitor of MAPKERK, were obtained from Calbiochem-Novabiochem Corporation (England, U.K.); SB-203580 and PD-98059 were reconstituted in dimethyl sulphoxide (DMSO), where the final concentration of DMSO was determined to be $\leq 0.01\%$. γ -Glutamylcysteinyl-ethyl ester (γ -GCE), a cell-permeant GSH analogue, was purchased from Calbiochem-Novabiochem Corporation and reconstituted in sterile deionized water and stored at 4°C. Phospho-MAPK^{p38} (Thr180/Tyr182) and MAPK^{p38} antibodies were purchased from New England Biolabs Incorporation (England, U.K.). Recombinant heat-shock protein 27 (Hsp27) was obtained from Calbiochem-Novabiochem Corporation and kept stored at -70° C.

Primary cultures of alveolar epithelia

Foetal alveolar type II (fATII) epithelial cells were isolated from lungs of foetuses, essentially as reported elsewhere (Haddad & Land, 2000a, b). Briefly, foetal rats were removed from pregnant Sprague-Dawley rats by caesarean section at day 19 of gestation (term = 22 days), the lungs excised, teased free from heart and upper airway tissue, and were finely minced then washed free of erythrocytes using sterile, chilled Mg²⁺- and Ca²⁺-free Hanks' balanced salt solution (HBSS; 0.5 ml foetus⁻¹). The cleaned lung tissue was resuspended in 1 ml foetus⁻¹ HBSS containing trypsin (0.1 mg ml⁻¹), collagenase (0.06 mg ml⁻¹) and DNase I (0.012% w v⁻¹), and was agitated at 37°C for 20 min. The solution was then centrifuged at $100 \times g$ for 2 min to remove undispersed tissue, the supernatant was saved to a fresh sterile tube and an equal volume of Dulbecco's modified Eagle medium (DMEM) with 10% (v v^{-1}) foetal calf serum (FCS) was added to the supernatant. After passing the supernatant through a 120 μ m pore sterile mesh, the filtrate was centrifuged at 420 $\times g$ for 5 min, the pellet re-suspended in 20 mls DMEM/FCS and the cells were placed into a T-150 culture flash for 1 h at 37°C to enable fibroblasts and nonepithelial cells to adhere. Unattached cells were washed three times by centrifugation at $420 \times g$ for 5 min each and then

seeded onto 24 mm diameter Transwell-clear permeable supports (Costar; 0.4 μ m pore size) at a density of 5×10⁶ cells per filter and were allowed to adhere overnight at 152 Torr (≈21% O₂/5% CO₂). DMEM/FCS was exchanged for 4 mls of serum free PC-1 media (Biowhittaker, MD, U.S.A.) pre-equilibrated to pO_2 =152 Torr and 37°C 24 h later and cells were maintained at this pO_2 until the experiment. In each case, and under conditions of independent treatment and pre-treatments, the adenylate energy charge, an index of cell viability and competence, remained ≥0.7 and transepithelial monolayer resistance was monitored constantly at 250–350 Ω cm² or more (Haddad & Land, 2000a; Haddad *et al.*, 2000).

Drug treatment and measurement of pro-inflammatory cytokine $TNF-\alpha$ by ELISA

Epithelial cells were pre-treated for 2 h with NAC (1-50 mM), washed twice in pre-equilibrated PC-1 medium and subsequently challenged with LPS (1 μ g ml⁻¹) for 24 h (LPS was derived from Escherichia coli, serotype 026: B6). Cell-free supernatants were assayed for pro-inflammatory (R&D Systems, U.K.) cytokine biosynthesis by two-site, solid phase, sandwich enzyme-linked immunosorbent assay (ELISA), essentially as recounted previously (Haddad et al., 2001a, c, d). Briefly, rabbit immunoaffinity purified polyclonal anti-rat TNF- α (2 µg ml⁻¹) antibody was used to coat high-binding microtitre plates (MaxiSorp, Nunc, U.K.) in bicarbonate buffer (0.1 M NaHCO3 and 0.1 M NaCl, pH 8.2) (Safieh-Garabedian et al., 1997); Haddad et al., 2001a, c). After blocking in 3% bovine serum albumin (BSA), recombinant (standard) and biotinylated (recognition) immunoaffinity purified sheep anti-rat cytokine antibodies were employed for secondary detection. The colour was developed using streptavidin-poly-HRP (Amersham Life Sciences, U.K.) coupled with 3,3',5,5'-tetramethyl-benzidine dihydrochloride (TMB) and 1 mM H₂O₂. The optical density was read at 450 nm against a background filter measuring at 595 nm, where the inter- and intra-assay coefficients of variations were reported at $\leq 10\%$. Results were extracted from the linear regression of the positive slope and cytokine concentration was expressed in pg ml^{-1} .

Measurement of intracellular levels of reduced (GSH) and oxidized (GSSG) glutathione

Reduced (GSH) glutathione concentrations were determined spectrophotometrically (Haddad & Land, 2000a; Haddad et al., 2000) in neutralized perchloric acid (PCA; 7%) extracts by following the glyoxylase-catalysed production of S-lactyl-GSH at 240 nm in a 1-ml volume containing 790 µl phosphate buffer (25 mM KH₂PO₄; 25 mM K₂HPO₄; pH 6.8), 150 µl 1% BSA, 10 µl sample, 10 µl glyoxylase-I (1 mg ml^{-1}) , and $40 \,\mu \text{l}$ methylglyoxal (0.1 M). Oxidized glutathione (GSSG) was determined in the same cuvette by addition of 1 mg ml^{-1} glutathione reductase and $8 \mu \text{l}$ of 12 mM β -NADPH and then by following the change in absorbance at 340 nm (Haddad & Land, 2000a; Haddad et al., 2000). Drift inherent to the assay was controlled by subtracting the absorbance change observed over the same time period from control cuvettes containing the same reaction components but with a matched sample volume of

deionized water. Protein content of each PCA precipitate was re-dissolved in 1 M NaOH and determined enabling results to be expressed as μ moles mg⁻¹ protein.

Preparation of subcellular extracts for Western analysis of p38 MAP kinase activation/phosphorylation

Analysis of p38/RK phosphorylation Total protein extraction was performed by homogenizing primary cell tissue in a suitable volume of a buffer $(1:40 \text{ w v}^{-1})$ containing 20 mM HEPES (pH 7.5), 1.5 mM MgCl₂, 0.2 mM EDTA and 0.1 M NaCl (Haddad & Land, 2000a; Haddad et al., 2000). Before extraction 5 mM dithiothreitol (DTT), 1 mM phenylmethylsulphonyl fluoride (PMSF), and 1.2 mM sodium orthovanadate (Na₃VO₄) were added to the buffer. The cellular debris was pelleted by centrifugation at $10,000 \times g$ for 30 min at 4°C, and the collected supernatant was mixed with an equal volume of the same extracting buffer but containing in addition 40% (v v⁻¹) glycerol. Threonine and tyrosine phosphorylation of p38 MAPK was analysed according to instructions given in commercially available kits (New England Biolabs, Inc., Beverly, MA, U.S.A.). The kit employs specific anti-phospho-p38 MAPK antibodies against Thr180/Tyr182 sites that do not cross-react with phosphorylated threonine/tyrosine of extracellular signal-regulated kinase (ERK) 1/2 or c-Jun-NH₂-terminal kinase (JNK). Analysis of Thr180/Tyr182 phosphorylation of p38 MAPK was performed as follows: Extracted proteins $(20-25 \mu g)$ were resolved over sodium dodecyl sulphate-polyacrylamide gel electrophoresis (SDS-PAGE; 7.5%) gels at RT, blotted onto nitrocellulose membrane, and non-specific binding sites were subsequently blocked. The membrane was probed with specific antibody to phosphorylated threonine and tyrosine of p38 MAPK for primary detection. Anti-rabbit Ig-biotinylated antibody (Amersham Life Science, U.K.) was employed for secondary detection followed by the addition of streptavidin-HRP conjugate and visualized on film by chemiluminescence. Phosphorylation-independent state of p38 MAPK using a specific antibody was used as an internal reference for semiquantitative loading in parallel lanes for each variable. Western blots were scanned by NIH MagiScanII and subsequently quantitated by UN-Scan-IT automated digitizing system (Version 5.1; 32-bit), and the ratio of the density of the band to that of the non-phosphorylated form was performed.

Analysis of Hsp27 phosphorylation by the upstream activated MAPKAP-K2 kinase Active phosphorylated p38 MAPK regulates a kinase cascade that may be followed by determining the terminal phosphorylation of Hsp27 by MAPKAP-K2. After treatment with LPS for the indicated doses and time points, cells were ruptured in 250 μ l of lysis buffer (mM): HEPES (pH 7.4) 20, EGTA 2, β -glycerophosphate 50, Na₃VO₄ 1, NaF 5, 1% (v v⁻¹) Triton X-100; 10% (v v⁻¹) glycerol; DTT 1, PMSF 1, 10 μ g/ml leupeptin; and 10 μ g/ml aprotinin, for 30 min on ice. Cell debris was removed by centrifugation at $10,000 \times g$ for 10 min at 4°C. The kinase activity of MAPKAPK-2, which is specifically regulated by the phosphorylation of the upstream p38 MAPK, was assayed with recombinant heat-shock protein 27 (Hsp27) as a substrate. Briefly, 1 μ g of Hsp27 was added to a microcentrifuge tube containing 10 μ l (\approx 10 μ g) of cellular extract. Kinase reaction was initiated by the addition of 10 μ l of γ^{-32} P-labelled Mg²⁺/ATP solution (10 mM MgCl₂/ 1 mM ATP/2 μ Ci of [γ^{-32} P]-ATP) and performed at 36°C for 20 min. The final concentration of Mg²⁺/ATP in the assay was adjusted to a 2.5 μ M and 0.25 μ M, respectively. Reactions were terminated by the addition of 5 μ l Laemmli sample buffer and subsequently boiled (95°C) for 5 min. Samples were separated by SDS-PAGE (17.5% (w v⁻¹) gel). Gels were blotted onto a Whatmann paper and dried for 2 h prior to exposure to autoradiography with a phosphorimager, followed by specific quantitation of the corresponding bands.

Selective inhibition of p38 MAP kinase and LPS-induced release of pro-inflammatory cytokines

Cells were pre-incubated for 2 h with SB-203580 (0, 0.01, 0.1, 1, 10, 100 μ M) (Calbiochem, U.K.), a selective inhibitor of MAPK p38/RK. This was followed by exposure to LPS (1 μ g ml⁻¹) for 24 h and cell-free supernatants were collected for cytokine analysis (TNF- α) by ELISA (Haddad *et al.*, 2001a, b, c), as recounted above.

Selective inhibition of p42/44 MAP ERK kinase and LPS-induced release of pro-inflammatory cytokines

Cells were pre-incubated for 2 h with PD-98059 (0, 0.01, 0.1, 1, 10, 100 μ M) (Calbiochem, U.K.), a selective inhibitor of MAPK p42/44/ERK. This was followed by exposure to LPS (1 μ g ml⁻¹) for 24 h and cell-free supernatants were collected for cytokine analysis (TNF- α) by ELISA (Haddad *et al.*, 2001a, b; c), as recounted above. Separately, cells were pre-treated (2 h) with a combination of SB-203580 and PD-98059, exposed to LPS (1 μ g ml⁻¹) for 24 h and subsequently assayed for TNF- α .

Selective inhibition of the nuclear factor- κB signalling pathway and the regulation of LPS-induced release of TNF- α

Cytokines have been demonstrated to induce NF- κ B, whose activation has been implicated in mediating biological responses, including the expression of genes encoding cytokines (Baldwin, 1996). Cells were pre-treated (2 h) with various inhibitors then monolayers washed twice in pre-equilibrated PC-1 medium and subsequently challenged with LPS $(1 \ \mu g \ ml^{-1})$ for 24 h. In order to determine whether LPSinduced release of TNF- α is regulated, at least in part, by NF- κ B, we used carbobenzoxy-L-leucyl-L-leucyl-L-leucinal (MG-132; Calbiochem, U.K.), a potent, reversible proteasome inhibitor $(K_i = 4 \text{ nM})$, reported to inhibit NF- κ B activation (Baldwin, 1996). Cells were pre-treated with MG-132 (0, 1, 10, 50 μ M) for 2 h before exposure to LPS for further 24 h. Sulfasalazine (SSA; 0, 0.1, 1, 10 mM) and caffeic acid phenylethyl ester (CAPE; 0, 1, 10, 100 μ M; Calbiochem, U.K.), which are potent specific inhibitors of NF- κ B, were also incorporated. TNF- α release was assayed (24 h) following challenge. Separately, cells were pre-treated for 2 h with SN-50 (0, 1, 10, 20 µM; Calbiochem, U.K.), a specific permeating inhibitor of NF- κ B nuclear translocation, followed by exposure to LPS (1 μ g ml⁻¹) for 24 h. This peptide contains the nuclear localization sequence (NLS) for the p50 NF- κ B subunit and the amino-terminal sequence of Kaposi fibroblast growth factor to promote cell permeability. An inactive mutant control for SN-50 peptide (SN-50 M; 20 μ M) corresponding to the same peptide sequence with substitutions of Lys³⁶³ for Asn and Arg³⁶⁴ for Gly in the NLS region has been used to confirm the specificity and selectivity of SN-50.

Assessment of intracellular reactive oxygen species (ROS) accumulation with LPS

For the determination of hydrogen peroxide (H2O2), cell supernatants post addition of LPS (1 μ g ml⁻¹) were collected and centrifuged at $2000 \times g$ for 5 min at 4°C, and then treated with 1 M NaOH. The change in absorbency at 600 nm was monitored against phenol red solution (PRS) containing (in mM): NaCl 140, K₃PO₄ 10, glucose 5.5, phenol red 0.28, and U ml⁻¹ horseradish peroxidase 20 (Haddad et al., 2001a, c). Standard curves using PRS and H2O2 were prepared (0-100 μ M), and results extrapolated from the linear regression were converted to nmoles/mg protein. For the quantification of superoxide anion (O_2^{-}) release, the supernatant was discarded after 24 h of culture, and cells were then reincubated with LPS in the presence of 80 µM ferricytochrome c (fc) suspended in HBSS. The amount of O_2^{-1} released was determined by measuring the absorgency at 550 nm against blanks containing fc and superoxide dismutase (SOD; 300 u ml⁻¹). All experiments were performed in duplicates and data are presented as the amount of O_2^{-1} released based on nmoles of reduced fc/min/mg protein (Haddad et al., 2001a, c). The relatively non-specific measure of 'OH production, which reacts with dihydrorhodamine (DHR), thereby yielding water and a tertiary rather stable free radical, allows this radical to undergo rearrangement of the π electrons, leading to formation of fluorescent rhodamine. Cells were cultured at 10⁵ well⁻¹ in flat-bottomed microtitre plates at normoxia. Wells were washed with sterile, pre-equilibrated saline, then culturing was continued in Kreb's/Ringer solution containing 50 µM DHR and LPS as indicated. Fluorescence was measured at excitation/emission wavelengths of 485/535 nm. Negative/positive controls were included to monitor the specificity of the Fenton reaction (Haddad et al., 2001a, c). The 'OH level measured of the control (no LPS) was calibrated to 100%, and variables were plotted against this level as logarithmic fluorescence units.

Exposure to ROS-generating systems and the role of antioxidants in $MAPK^{p38}$ *signalling*

The rate of induced production of $O_2^{-\cdot}$ by xanthine (X) in the presence of xanthine oxidase (XO) was measured as the reduction of fc, as detailed previously (Haddad *et al.*, 2001a, c). X (100 μ M) and XO (0–2 mU ml⁻¹) were incubated in microtiter plates coated with 10⁵ cells well⁻¹ with 100 μ l of reaction solution containing 50 μ M fc in HBSS. The degree of fc reduction was measured at 550 nm. The standardized reaction allows production of O₂⁻ at an average rate of $\approx 8.10 \pm 0.09$ nmoles min⁻¹ ml⁻¹ (XO=2 mu ml⁻¹). Separately, cells were incubated in the presence of X (100 μ M) and XO (2 mu ml⁻¹) or H₂O₂ (100 μ M) for 15 min at 37°C. Subcellular extracts were prepared for assessing MAPK^{p38} phosphorylation/activation as recounted above. Independently, in order to investigate the possibility of ROS acting as signalling messengers in MAPK^{p38} signalling, selective antioxidants against O2-, OH and H2O2 radicals were employed. This choice was based on the antioxidant potency and specificity: Dimethyl sulphoxide (DMSO) is a prototypical scavenger of 'OH (Parker et al., 1985; Wasil et al., 1987, 1,3-dimethyl-2-thiourea (DMTU) scavenges H₂O₂ and 'OH (Fox, 1984; Parker et al., 1985; Wasil et al., 1987), and 4'-hydroxy-3'-methoxy-acetophenone (HMAP) is an NADPH-oxidase inhibitor which may also block ROS production by mitochondria (Lapperre et al., 1999). Epithelial cells were pre-treated for 2 h with DMSO (1%), DMTU (10 mM) and HMAP (10 mM), before exposure to LPS (1 μ g ml⁻¹) for 15 min. Extracts were then prepared for the assessment of MAPK^{p38} phosphorylation/activation. Separately, in order to determine the level of participation of mitochondrial ROS, cells were pre-treated with diphenylene iodonium (DPI; 10 µM), a non-specific inhibitor of flavin-dependent O2- producing enzymes, including not only mitochondrial but also the classical NADPH oxidase (Chandel et al., 1998), or KCN (10 mM), an inhibitor of mitochondrial respiratory chain, followed by exposure to LPS (1 μ g ml⁻¹) for 15 min. In addition, γ -GCE (100 μ M), a cell-permeant precursor of GSH, was added to monolayers 2 h before exposure to LPS (1 μ g ml⁻¹) for 15 min, followed by assessment of MAPK^{p38} phosphorylation/activation.

Statistical analysis and data presentation

Data are the means and the error bars the s.e.mean. Statistical evaluation of the difference in mean separation was performed by one-way analysis of variance (ANOVA), followed by *post hoc* Tukey's test, and the *a priori* level of significance at 95% confidence level was considered at $P \leq 0.05$.

Results

The excitatory role of LPS in mediating the phosphorylation of MAPK p38/RK

The induction of threonine/tyrosine phosphorylation of p38/ RK MAP kinase reflects the activation state of this kinase in regulating the downstream pathway involving MAPKAP-K2. In addition, the role of LPS in mediating the phosphorylation of p38/RK MAPK is not well characterized in the foetal alveolar epithelium. Consequently, we examined the phosphorylation of this kinase in response to LPS in controlled time- and dose-dependent experiments. As shown in Figure 1A, exposure of alveolar epithelial cells to LPS (1 μ g ml⁻¹) induced, in a time-dependent manner, the phosphorylation of p38/RK kinase (p-p38) on threonine/tyrosine residues as determined by a selective antibody probed on an SDS-PAGE analysis gel (see Methods). The positive control (+ve control) contained protein extracts prepared from C-6 glioma cells stimulated with anisomycin to induce the phosphorylation of threonine and tyrosine residues of p38/RK kinase. The phosphorylation state-independent p38/RK MAP kinase is shown in the lower panel of Figure 1A to verify semiquantitative loading for gel analysis per loading lane. The amount of p38/RK phosphorylation was quantified by an image analyser to indicate that the peak of p38/RK phosphorylation due to LPS stimulation is somewhere

around the 15-min time point, as shown in the histogram (Figure 1B). This reference time point (15 min) was subsequently adopted in further experiments. The phosphorylation of p38/RK MAP kinase became immediately active within minutes post addition of LPS (significantly active at 2 min), continued to increase up until 30 min, thereafter declining, still significantly different at 60 min, but its activity was lost between 90-120 min (Figure 1A,B). Figure 1C shows the dose-dependent analysis of LPS stimulation (0-1000 ng ml⁻¹) at 15 min, indicating a maximum induction of p38/RK phosphorylation at 100-1000 ng ml⁻¹ concentration range (Figure 1D). The steady, phosphorylation-independent (p38) was also quantitated at various doses of LPS (Figure 1C) to account for equal loading per lane. The specificity of SB-203580 was confirmed by the fact that this compound selectively prevented LPS-mediated phosphorylation of MAPK^{p38} without affecting the phosphorylation-dependent state of either JNK or ERK kinase (Data not shown).

The effect of LPS on the activation of MAPKAP-K2, the downstream kinase controlled by p38/RK MAP kinase, as assessed by the phosphorylation of its substrate, Hsp27

The activity of MAPKAP-K2 was evaluated by the phosphorylation state of its substrate, Hsp27. A representative SDS-PAGE Western gel showing the effect of LPS (1 μ g ml⁻¹) on the amount of γ -³²[ATP] bound to Hsp27 indicating the degree of its phosphorylation (p-Hsp27) is shown in Figure 2A, exhibiting the time response curve. As shown in Figure 2A, the peak time for Hsp27 phosphorylation consistently occurred at a later time (30-60 min) than the activation of p38/RK MAP kinase (15 min). The time peak was determined by histogram analysis of the phosphorylated band, as shown in Figure 2B. The dose-response curve showing the effect of LPS at 45 min on Hsp27 phosphorylation is shown in Figure 2C. There was gradual increase in the activity of MAPKAP-K2 as assessed by Hsp27 phosphorylation on LPS exposure, peaking at between 10-100 ng ml⁻¹, thereafter declining but still statistically significant at 10,000 ng ml⁻¹ (Figure 2D).

The role of p38/RK signalling pathway and

N-acetyl-L-cysteine (*NAC*) pre-treatment in mediating the effect of LPS on TNF- α biosynthesis

The activity, but not the activation, of p38/RK MAP kinase is selectively blocked by the pyridinyl imidazole compound SB-203580, thereby blockading the downstream pathways associated with the activation of MAPKAP-K2, Hsp27 phosphorylation and pro-inflammatory cytokine transcription and biosynthesis (Garrington & Johnson, 1999). As shown in Figure 3A, exposure of alveolar epithelial cells to medium alone for 24 h had no effect on the release of TNF- α into the supernatant. In addition, pre-treatment with SB-203580 (100 μ M) for 1 h and further exposure to medium alone had no apparent effect on TNF-a secretion. In contrast, administration of LPS (1 μ g ml⁻¹) for 24 h enhanced the secretion of TNF- α into the supernatant (Figure 3A). Pretreatment with SB-203580, followed by exposure to LPS blockaded, in a dose-dependent manner, LPS-dependent biosynthesis of TNF- α , as shown in Figure 3A. At all concentrations of SB-203580 used, there were no signs of cytotoxicity associated with the treatment, nor were there any



Figure 1 The immunoregulatory role of lipopolysaccharide (LPS) in augmenting the phosphorylation and activation of p38 MAPK. Foetal alveolar epithelial cells were stimulated with LPS (1 μ g ml⁻¹) for the desired time points as indicated. Lysates were separated by a 7.5% SDS-PAGE, transferred to membranes and probed with specific antibodies recognizing the phosphorylated and steady state forms of p38 MAPK. (A) LPS induced, in a time-dependent manner, the phosphorylation of p38, with maximum induction at between 10–15 min (*p*-p38 indicates the phosphorylated form and p38 that of the steady-state form). The positive control lane contained loading of anisomycin extract for the recognition of the specificity of phosphorylated p38 bands. (B) Histogram analysis of p38 phosphorylation for the time-response curve and *Pseudo-Voight* curve fitting showing the effect of LPS (15 min) on p38 phosphorylation and steady-state forms, with maximum induction at a concentration of 1000 ng ml⁻¹. (D) Histogram analysis of p38 phosphorylation and maximum induction with LPS. **P*<0.05, ***P*<0.01, *** *P*<0.001, as compared with control (LPS=0 ng ml⁻¹). *n*=3, which represents the number of independent experiments with separate cell preparations.

decreases in the transepithelial resistance of monolayers (Data not shown). Figure 3B shows the dose-dependent inhibition of LPS-induced TNF- α biosynthesis due to pretreatment (2 h) with *N*-acetyl-L-cysteine (NAC), an antioxidant and a precursor of cysteine, the rate-limiting amino acid in the biosynthesis of glutathione (GSH) (Meister, 1988; Rahman, 1999; Haddad & Land, 2000a; Haddad *et al.*, 2000).

The role of p42/44/ERK signalling pathway and its synergism with p38/RK in mediating the effect of LPS on TNF- α biosynthesis

The effect of PD-98059, a selective inhibitor of MAPK^{ERK} pathway, on LPS-mediated TNF- α secretion is shown in Figure 4A. Exposure of alveolar epithelial cells to medium alone for 24 h had no effect on the release of TNF- α into the supernatant. In addition, pre-treatment with PD-98059 (100 μ M) for 1 h and further exposure to medium alone had no apparent effect on TNF- α secretion. In contrast, administration of LPS (1 μ g ml⁻¹) for 24 h enhanced the secretion of TNF- α into the supernatant (Figure 4A). Pre-treatment with PD-98059, followed by exposure to LPS blockaded, in a dose-dependent

manner, LPS-dependent biosynthesis of TNF- α , but to a lesser extent than SB-203580 (Figures 3A, 4A). At all concentrations of PD-98059 used, there were no signs of cytotoxicity associated with the treatment, nor was there any decrease in the transepithelial resistance of monolayers (Data not shown). A combination of SB-203580 and PD-98059 exhibited a mild synergistic effect, especially at a dose of 0.01 μ M, which was ineffective with either inhibitor used on its own (Figure 4B).

The effect of proteasome inhibitor and selective inhibition of NF- κ B in mediating the effect of LPS on TNF- α biosynthesis

As shown in Figure 5A, exposure of epithelial cells to medium alone for 24 h had no effect on the secretion of TNF- α . Similarly, pre-treatment of monolayers with MG-132 (50 μ M), a specific inhibitor of the proteasome complex, reportedly known to blockade the NF- κ B pathway, followed by exposure to medium alone for 24 h had no effect on TNF- α production either (Figure 5A). Exposure to LPS (1 μ g ml⁻¹) for 24 h induced TNF- α biosynthesis $\approx 3-4$ fold relative to medium alone measured at the same time point. Pre-treatment with MG-132 prior to



Figure 2 The regulation by LPS of the activity of MAPKAP-K2, the downstream kinase activated by p38 phosphorylation, through the phosphorylation of its substrate, Hsp27. (A) LPS induced, in a time-dependent manner, the phosphorylation of Hsp27 (p-Hsp27), an index of the activity of MAPKAP-K2, with peak analysis at 30-60 min. (B) Histogram analysis of Hsp27 phosphorylation for the time-response curve and Pseudo-Voigt curve fitting showing the peak at ≈45 min. (C) Dose-response curve showing the effect of LPS (45 min) on Hsp27 phosphorylation, with maximum induction at a concentration of 1000 ng ml⁻¹, thereafter declining. (D) Histogram analysis of Hsp27 phosphorylation for the dose-response curve and Pseudo-Voigt curve fitting showing gradual and maximum induction with LPS. *P<0.05, **P<0.01, ***P<0.001, as compared with control $(LPS = 0 \text{ ng ml}^{-1})$. n = 3, which represents the number of independent experiments with separate cell preparations.



Figure 3 The effect of selective inhibition of p38 MAPK and NAC pre-treatment on LPS-induced TNF-a biosynthesis. (A) Pretreatment with SB-203580 (2 h), a selective inhibitor of p38 MAPK, prior to exposure to LPS (1 µg ml⁻¹), reduced LPS-induced TNF- α production in a dose-dependent manner. LPS upregulated the secretion of TNF- α approximately 4 fold relative to cells incubated in medium alone (control). Pre-treatment with SB-203580 (100 μM) on its own did not affect TNF-α synthesis. SB-203580 reduced LPS-induced TNF- α secretion at doses $\ge 0.1 \ \mu$ M, with maximum inhibition at 100 μ M (IC₅₀=0.054 \pm 0.005 μ M). (B) Pretreatment with NAC (2 h) before stimulating with LPS reduced, in a dose-dependent manner, the induction of $TNF-\alpha$, effective at doses ≥ 1 mM, with maximum suppression at 50 mM (IC₅₀=26.07±2.25 mM). $\phi P < 0.05$, as compared with control; *P < 0.05, **P < 0.01, ***P < 0.001, as compared with LPS (1 μ g ml⁻¹). n = 4, which represents the number of independent experiments run in duplicate with separate cell preparations.



Figure 4 The effect of selective inhibition of p42/44 MAPK on LPS-induced TNF- α biosynthesis. (A) Pre-treatment with PD-98059 (2 h), a selective inhibitor of MAPK^{ERK}, prior to exposure to LPS (1 μ g ml⁻¹), reduced LPS-induced TNF- α production in a dose-dependent manner. LPS upregulated the secretion of TNF- α approximately 4 fold relative to cells incubated in medium alone (control). Pre-treatment with PD-98059 (100 μ M) on its own did not affect TNF- α synthesis. PD-98059 reduced LPS-induced TNF- α secretion at doses $\ge 0.1 \ \mu$ M, with maximum inhibition at 100 μ M (IC₅₀=5.12±0.47 μ M). (B) Pre-treatment with a combination of SB-203580 and PD-98059 synergistically reduced, in a dose-dependent manner, the induction of TNF- α , effective at doses $\ge 0.01 \ \mu$ M, with maximum and complete suppression at 100 μ M (IC₅₀=0.023±0.003 μ M). $\phi P < 0.05$, as compared with control; *P < 0.05, **P < 0.01, ***P < 0.001, as compared with LPS (1 μ g ml⁻¹). n = 3, which represents the number of independent experiments run in duplicate with separate cell preparations.

challenge with LPS did not affect or reduce LPS-dependent release of TNF- α (Figure 5A). As shown in Figure 5B, pretreatment of monolayers with CAPE (100 μ M), a specific inhibitor of NF- κ B, followed by exposure to medium alone for 24 h had no effect on TNF-α production either (Figure 5B). Pretreatment with CAPE prior to challenge with LPS reduced LPSdependent release of TNF-a at a minimum effective dose $\geq 10 \ \mu M$ (Figure 5B). Figure 5C shows that pre-treatment of monolayers with SSA (10 mM), a potent and specific inhibitor of NF- κ B, followed by exposure to medium alone for 24 h had no effect on TNF- α production either (Figure 5C). Pre-treatment with SSA prior to challenge with LPS reduced LPS-dependent release of TNF- α at the highest dose used in this study (10 mM) (Figure 5C). As shown in Figure 5D, pre-treatment of monolayers with SN-50 (20 µM), a potent and specific permeating inhibitor of NF- κ B, followed by exposure to medium alone for 24 h had no effect on TNF- α production either (Figure 5D. Pre-treatment with SN-50 prior to challenge with LPS reduced, in a dose-dependent manner, LPS-dependent release of TNF- α (Figure 5D). Mutation of the SN-50 peptide (SN-50M; 20 μ M) reversed the inhibitory effect of this peptide on LPS-mediated TNF-α biosynthesis (Figure 5D). The 50% minimum inhibitory effective concentration (IC50) of NF-kB inhibitors on LPSmediated TNF- α biosynthesis is given in Table 1.

The effect of LPS on intracellular accumulation of ROS and the role of antioxidants and NAC

The underlying mechanism of the mode of action of LPS as a mediator of oxidative stress was investigated. LPS induced, in a time-dependent manner, the accumulation of H_2O_2 (Figure 6A), O_2^{--} (Figure 6B) and OH (Figure 6C). In order to assess the

likely source of ROS in response to LPS, cells were pre-treated with HMAP or NAC prior to exposure to LPS for 30 min. HMAP and NAC reduced H_2O_2 accumulation (Figure 6D), but to a lesser extent than O_2^{-1} (Figure 6E). The effect of HMAP on 'OH was milder than that of NAC (Figure 6F). This indicates that the membrane-bound NADPH oxidase and the mitochondrial complex are likely involved as potential resources for ROS in response to LPS, an effect which is redox-sensitive.

The effect of ROS-generating systems and antioxidants on LPS-mediated $MAPK^{p38}$ phosphorylation

Exposure to O_2^{-1} -generating system (X/XO; 100 μ M/ 2 mu ml⁻¹) induced the phosphorylation of MAPK^{p38}, as shown in Figure 7A. Similarly, H₂O₂ (100 μ M), which is a major source of OH, up-regulated the phosphorylation of MAPK^{p38}, as shown in Figure 7A. Pre-treatment with selective antioxidants, purported to reduce the intracellular accumulation of H₂O₂, O₂⁻¹ and OH, abrogated LPS-mediated phosphorylation of MAPK^{p38}, as shown in Figure 7B. The antioxidant and cellpermeant precursor of GSH, γ -GCE, reduced MAPK^{p38} phosphorylation (Figure 7B). The numbers shown below the upper panel of Figure 7B exhibit the per cent phosphorylation state of MAPK^{p38}. The lower panels in Figure 7A,B show the steady state, phosphorylation-independent, MAPK^{p38}.

The role of NAC in mediating glutathione (GSH) biosynthesis

The cysteine that is provided due to NAC administration is eventually fed into the biosynthetic pathway that brings about the formation of GSH by the action of γ -glutamylcysteine synthetase (γ -GCS), the rate-limiting enzyme in the biosynthesis of GSH (Meister, 1988; Hayes & McLellan, 1999; Haddad & Land, 2000a). Incubation of alveolar

epithelial cells for 24 h with NAC induced, in a dosedependent manner, the intracellular accumulation of GSH (Figure 8A). This effect was associated with a dose-dependent



Figure 5 The role of the NF-κB signalling pathway in regulating LPS-mediated TNF-α biosynthesis. (A) Incubation of epithelial cells with medium alone (24 h) had no effect on TNF-α production, compared to pre-incubation with MG-132 (2 h), a proteasome inhibitor, followed by exposure to medium alone (24 h). Exposure to LPS (1 µg ml⁻¹; 24 h) induced TNF-α secretion into the supernatant, an effect that was not affected or reversed by MG-132, at all doses tested. (B) The effect of CAPE on LPS-mediated TNF-α biosynthesis, where incubation with medium alone had no effect on TNF-α production, compared to pre-incubation with CAPE (2 h), an inhibitor of NF-κB, followed by exposure to medium alone. The effect of LPS was reversed by CAPE at doses ≥ 10 µM. (C) The effect of SSA on LPS-mediated TNF-α biosynthesis, where pre-incubation with CAPE, followed by exposure to medium alone had no effect. The effect of LPS was reversed by SSA only at a dose of 10 mM. (D) The effect of SN-50 and its mutant, SN-50M, on LPS-mediated TNF-α biosynthesis showed that pre-incubation with SN-50, followed by exposure to medium alone had no effect on TNF-α. The effect of LPS was reversed by SN-50 in a dose-dependent manner. Mutation of the wild-type peptide (SN-50 M; 20 µM) reversed the inhibitory effect of SN-50 on LPS-dependent TNF-α secretion. $\phi P < 0.05$, as compared with LPS alone. n=3, which represents the number of independent experiments run in duplicate.



Figure 6 Analysis of ROS accumulation in response to LPS and the role of selective antioxidants. (A) H_2O_2 began to increase at 15 min and continued to be significantly higher than control until 24 h, thereafter declining. (B) O_2^{-} appeared at 15 min and continued to be higher than control until 24 h, thereafter declining. (C) The concentration of OH (fluorescence) began to increase at 15 min and continued to be significantly higher than control until 24 h, thereafter declining. (D) HMAP and NAC pre-treatments reduced accumulation of H_2O_2 in response to LPS. (E) HMAP and NAC reduced accumulation of O_2^{-} in response to LPS, in a dose-dependent manner. (F) HMAP and NAC mildly reduced accumulation of 'OH in response to LPS. $\phi P < 0.05$, as compared with control; *P < 0.05, **P < 0.01 ***P < 0.001, as compared with control in the absence or presence of LPS. n=4, which represents the number of independent experiments run in duplicate.

Table 1 Analysis of the 50% minimum effective inhibitory (IC₅₀) concentrations of various NF- α B inhibitors on LPS-induced TNF- α biosynthesis in the alveolar epithelium

MG-132 (µм)	$CAPE~(\mu M)$	SSA (mm)	SN-50 (µм)
-	93.15 ± 5.42	≥10	21.86 ± 2.57

Data are presented as mean \pm s.e.mean. n = 3, each. IC₅₀ was determined from the negative slope of the linear regression curves.

decrease in the intracellular level of the oxidized disulphide gultathione (GSSG), as reflected by an increasing redox ratio of GSH/GSSG (Figure 8A). The time-response curve for NAC incubation at 50 mM is shown in Figure 8B. Incubating cells with NAC for different time points induced the intracellular formation of GSH and reduced the levels of GSSG (Figure 8B). Within 2 h of NAC incubation there was significant accumulation of GSH, the concentration of which continued to elevate in the continuiing presence of NAC and maximized at around 16–24 h, thereafter declining (48 h) but persistently found to be still significantly different from control deproteinated monolayers (Figure 8B). Redox equilibrium assessment of glutathione ratio homeostasis reported in alveolar cells treated with NAC for 24 h at 37° C is given in Table 2.

The role of NAC in regulating the $MAPK^{p38}$ signalling pathway

The amount of threonine/tyrosine phosphorylation of p38/ RK MAP kinase was reduced, in a dose-dependent manner, with NAC pre-treatment (2 h) prior to exposure to LPS (1 μ g ml⁻¹) for 15 min (Figure 9A). LPS-mediated phosphorylation of p38/RK MAP kinase was maximally blockaded with NAC at 50 mM. The steady, phosphorylation-independent state of MAPK^{p38} is shown in the lower panel of Figure 9A. The dose-response curve exhibiting the inhibitory effect of NAC on MAPK^{p38} phosphorylation state is given in a form of histogram analysis in Figure 9B.

Discussion

TNF- α is a potent inflammatory cytokine, which exerts its pleiotropic activities through ligand-induced cross-linking of specific receptors, virtually present in almost all cell types



Figure 7 The effect of ROS-generating systems and selective antioxidants on MAPK^{p38} phosphorylation. (A) Xanthine (X)/ xanthine oxidase (XO) system, which generates O_2^{-} , up-regulated the phosphorylation of MAPK^{p38}, similar to the effect of exogenous H₂O₂, which is a major source for OH. (B) Selective antioxidants, including DMSO, DMTU, HMAP, DPI and KCN, attenuated LPS-dependent MAPK^{p38} phosphorylation. The antioxidant and cell-permeant precursor of GSH, γ -GCE, similarly attenuated the effect of LPS (*p*-p38 indicates the phosphorylated form and p38 that of the steady-state form). *n*=3, which represents the number of independent experiments.

(Vandenabeele *et al.*, 1995). During inflammation, TNF- α released in the inflammatory environment transiently activates neutrophils and macrophages, thereby causing them to release O_2^{-1} as a consequence of the activation of the plasma membrane NADPH oxidase (Klebanoff *et al.*, 1986). This oxidative burst features a rapid but transient release of ROS and is a crucial part of the defence mechanism against invading pathogens and tumour cell metastasis. Although TNF- α is primarily produced by macrophages, growing evidence suggested that other cell types, such as endothelial and epithelial cells, release TNF- α and other inflammatory mediators, thereby amplifying and boosting the inflammatory cells (Thompson *et al.*, 1985; Hashimoto *et al.*, 1999b; Haddad *et al.*, 2001a, c, d).

There is increasing evidence implicating the alveolar epithelium, in particular, as a dynamic barrier that plays an important role in regulating the inflammatory and metabolic responses to oxidative stress and the accompanying inflammatory signal, sepsis, endotoxaemia, and other critical illnesses in the lung (Thompson *et al.*, 1985; Pittet *et al.*, 1997; Haddad *et al.*, 2001a, b, c, d). The respiratory epithelium is a primary target of an inflammatory/infectious condition at the epithelial-blood interface and is itself capable of amplifying an inflammatory signal by recruiting inflammatory cells and by producing inflammatory mediators. Many of the side effects of LPS, derived from the cell wall of gram-negative bacteria, are secondary to the overproduction of pro-inflammatory mediators. Inflammatory as well as autoimmune disease is often associated with deregulated expression and biosynthesis of pro-inflammatory cytokines, including TNF- α , which influence a plethora of cellular functions. Therefore, the down-regulation of an inflammatory signal is a major focus of the rational approach to the treatment of inflammatory diseases, such as chronic inflammation, sepsis and rheumatoid arthritis. For instance, a novel recent study by Haskó et al. (2000) reported a potential role for extracellular purines, including adenosine and ATP, and inosine, a degradation product of these purines, as potent endogenous immunomodulatory molecules that inhibit inflammatory cytokine biosynthesis and protect against endotoxin-induced shock. It has also been reported, in addition, that selective inhibition of phosphodiesterases, a family of enzymes involved in the degradation of cyclic AMP (Haskó et al., 1998; Doherty, 1999), steroids, such as glucocorticoids (Visser et al., 1998), pyrimidylpiperazine derivatives (Fukuda et al., 2000; Hanano et al., 2000; Hisadome et al., 2000; Haddad et al., 2001e), and ERK and p38/RK MAPK selective inhibitors (Su & Karin, 1996; Garrington & Johnson, 1999; Haddad, 2001) differentially regulate the transcription and biosynthesis of inflammatory cytokines.

The novel mammalian MAPK^{p38} was originally identified in murine pre-B lymphocytes transfected with the LPScomplex receptor CD14 and in murine macrophages where it



Figure 8 The effect of NAC on intracellular redox potential (GSH/ GSSG). (A) Cells were exposed to NAC treatment for 24 h, thereby inducing the intracellular accumulation of the reduced form of glutathione (GSH) in a dose-dependent manner. Intracellular accumulation of GSH due to NAC was at the expense of the oxidized form of glutathione (GSSG), as evident from the reduced GSSG/GSH ratio. (B) Time-dependent analysis of the effect of NAC (50 mM) on GSH and GSSG accumulation. GSH concentration prevailed with NAC treatment, statistically significant as early as 2 h post incubation with NAC, thereafter ascending to a maxima at 24 h, declining, but still persistently significant, at 48 h. This elevation of intracellular GSH due to NAC is partly at the expense of GSSG, as shown from the reduced GSSG/GSH equilibrium ratio. $\phi P < 0.05$, $\phi\phi P < 0.01$, $\phi\phi\phi P < 0.001$, for (GSH) as compared with (GSSG); *P < 0.05, **P < 0.01, ***P < 0.001, as compared with control (NAC at 24 h and NAC at 50 mM); +P < 0.05, ++P < 0.01, +++P < 0.001, for (GSSG) as compared with (GSSG)_{control}.

Table 2	Redox	equilibr	ium ass	essmen	it of gl	utathi	one r	atio
homeosta	isis repo	rted in	alveolar	cells t	reated	with	NAC	for
24 h at 3	7°C ¯							

Glutathione ratio ((GSH)+(GSSG))/((GSSG))

Degree of reduction equilibrium with NAC (mm)

0	1	10	50
2.51 ± 0.15	3.76 ± 0.17	$5.97 \pm 0.65*$	$7.55 \pm 0.78 **$

Data are means \pm s.e.mean. NAC (0; n = 5), (1 mM; n = 5), (10 mM; n = 5) and (50 mM; n = 4). *P < 0.05, **P < 0.01, ***P < 0.001, as compared to control (0), without NAC treatment. *n* refers to number of measurements taken from at least three independent cell filters, where the entire results were pooled and averaged.

was activated in response to LPS (Han et al., 1994). In parallel, MAPK^{p38} was also identified as a reactivating kinase (RK), which activates MAPKAP kinase-2 (MAPKAP-K2; MK-2), which in turn phosphorylates the small heat-shock protein, Hsp27, and regulates the stability of cytokine transcripts bearing the pentanucleotide sequence of AU-rich elements (ARE) (Rouse et al., 1994; Su & Karin, 1996; Rutault et al., 2001). Of note, MAPK^{P38}, MAPK^{ERK} and the downstream pathways regulated are essentially crucial for LPS-induced cytokine gene expression and biosynthesis (Lee et al., 1994; Cano & Mahadevan, 1995; Su & Karin, 1996; Garrington & Johnson, 1999). For instance, knockout mice lacking MK-2 (MK- $2^{-/-}$) exhibited a colossal reduction in the biosynthesis of TNF- α , a potent pro-inflammatory cytokine that is involved in many human diseases (Hudson et al., 1995; Pittet et al., 1997; Herlaar & Brown, 1999), and are remarkably resistant to shock induced by LPS (Kotlyarov et al., 1999). Recent evidence, furthermore, suggested that the MAPK^{p38}-MAPK/MK-2 pathway is redox-sensitive and governs the transcription and biosynthesis of pro-inflammatory cytokines in a ROS and redox-dependent mechanism (De Keulenaer et al., 2000; Kang et al., 2000; Chan et al., 2001; Hashimoto et al., 2001). However, redox/ROS regulation of the MAPK^{p38/ERK} -MAPK/MK-2 pathway mediating TNF- α signalling in the foetal alveolar epithelium is not well characterized. It was subsequently hypothesized that MAPK^{p38/ERK}-mediated regulation of LPS-dependent TNF- α biosynthesis is redox-sensitive and that intracellular glutathione manipulation modulates the capacity of the epithelium to produce this inflammatory cytokine.

Redox regulation of MAPK-mediated TNF-*α* biosynthesis

The signal transduction cascade that regulates the biosynthesis and secretion of TNF- α in the foetal alveolar epithelium has not been well defined. Administration of LPS, derived from E. coli, up-regulated the extracellular accumulation of TNF- α and other cytokines (Haddad *et al.*, 2001a,c,d). The effects of LPS are essentially mediated by the membrane-bound CD14 complex and, in part, through the circulating LPS-binding protein (LBP) (Wright et al., 1990; Rietschel et al., 1994). It was reported, furthermore, that LPS-mediated responses on TNF-a transcription and biosynthesis bifurcate at the level of Ras/Raf G-coupled proteins into two major signalling pathways: One that runs through NF- κ B-inducing kinase (NIK) route, which regulates the phosphorylation of the inhibitory- κB (I κB) proteins, the cytosolic inhibitors of NF- κ B, and another which is mediated through the extracellular signal-regulated kinase (MAPK^{ERK}) and MAPK^{p38} pathways (Su & Karin, 1996; Garrington & Johnson, 1999; Mercurio and Manning, 1999). The promoters of genes encoding cytokines contain multiple cis-acting motifs including those that bind such transcription factors as NF- κ B. Furthermore, the release of free NF- κ B upon extracellular stimulation due to IkB phosphorylation and degradation by the proteasome complex, leads to DNA binding to specific κB moieties in order to initiate transcription of related genes, including immunoreceptors, cytokines and, interestingly, its own inhibitor, IkB (Baldwin, 1996; Haddad et al., 2001d, e). Two unique features of the NF- κ B/I κ B complex system are deduced from its feedback regulation. The transcriptional activation of NF-kB triggers the synthesis of $I\kappa B$, and NF- κB activated transcription is maintained by continuous degradation of $I\kappa B$, which is sustained by an extracellular stimulus. Thus, the expression



Figure 9 The attenuating effect of NAC on p38/RK phosphorylation. (A) Epithelial cells were cultured with medium alone (control), pre-treated with NAC (50 mM) for 2 h prior to exposure to medium alone for 15 min, exposed to LPS (1 μ g ml⁻¹) for 15 min, or pre-treated with NAC (1-50 mM) for 2 h followed by exposure to LPS (1 μ g ml⁻¹) for 15 min. Lysates were separated by a 7.5% SDS-PAGE and probed with specific antibodies directed against the phosphorylated threonine/tyrosine residues of p38/ RK MAP kinase or the phosphorylation-independent state of p38. Pre-treatment with NAC (50 mM) followed by exposure to medium alone did not affect the phosphorylation of p38/RK, despite the observation it abolished the faint band appearing under the control lane. Pre-treatment with NAC attenuated the phosphorylation-mediated effect of LPS, in a dose-dependent manner, completely blocking its phosphorylation at 50 mM (IC₅₀=20.62 \pm 3.08 mM). The steady state of p38 is largely not affected with any of the aforementioned treatments. (B) Histogram analysis of p38 phosphorylation for the inhibitory dose-response curve of NAC. $\phi P < 0.05$, as compared with control; *P < 0.05, **P < 0.01, ***P < 0.001, as compared with control (LPS = 1 μ g ml⁻¹). n = 3, which represents the number of independent experiments with separate cell preparations.

of I κ B parallels both NF- κ B activity and the duration of the activating extracellular stimulation, suggesting that this temporal parallelism between IkB accumulation/degradation and an effective external stimulation is a mechanism allowing dual regulation of NF- κ B within the alveolar space. The selective interference with the functioning of the proteasome complex in regulating the translocation and activation of NF- κB and the expression of its inhibitor I κB - α suggested that the $I\kappa B - \alpha/NF - \kappa B$ pathway is partially implicated in regulating LPS-mediated biosynthesis of TNF- α . This is rather unequivocally reinforced with the observation reported in this study that selective inhibition of the NF- κ B pathway attenuated but did not abrogate the inductive effect of LPS on TNF- α production. Therefore, the I κ B- α /NF- κ B pathway could be partially dissociated from the presumably downstream pathway regulating TNF- α signalling, indicating the involvement of a possible cross-talk among several pathways, such as MAPK^{p38/ERK}, working independently or in coherence to integrate signalling mechanisms governing the regulation of cytokines in the alveolar epithelium (Nemoto et al., 1998).

Exposure to LPS induced, in a time- and dose-dependent manner, the phosphorylation of MAPK^{p38}, suggesting the involvement of an up-stream kinase, such as MAPK kinase (MKK), in LPS-mediated regulation of the MAPK^{p38} pathway (Widmann et al., 1999). Similar to other MAPK families, MAPK^{p38} is activated by dual phosphorylation of threonine (Thr) and tyrosine (Tyr) in the so-called Thr-Gly-Tyr activation loop or motif (Su & Karin, 1996; Garrington & Johnson, 1999; Widmann et al., 1999). Upon stimulation, MAPK^{p38} phosphorylates and activates the MK-2 pathway, which regulates the phosphorylation of Hsp27. The biological activity of MAPK^{p38} is selectively blocked by the pyridinylimidazole compound SmithKline Beecham ((SB-203580), which is postulated to possess anti-inflammatory activity (Widmann et al., 1999). In this respect, SB-203580 has been reportedly associated with the suppression and/or augmentation of the transcription and biosynthesis of a wide array of inflammatory and anti-inflammatory mediators, including IL-1β (Baldassare et al., 1999), IL-6, IL-8 (Griego et al., 2000; Hashimoto et al., 2001), IL-10 (Foey et al., 1998; De et al., 2000), IL-12 (Widmann et al., 1999), TNF-α (Yamakawa et al., 1999; Rutault et al., 2001) and prostaglandin-E2 (PGE2) (Scherle et al., 1998). Our results jibe with the proposal that selective blockade of the MAPK^{p38} signalling transduction pathway exerts an anti-inflammatory condition via suppression of the release of inflammatory mediators, including TNF- α . Despite the fact that selective blockade of the MAPKERK pathway partially attenuated LPS-dependent secretion of TNF-a, simultaneous inhibition of MAPK^{p38} and MAPKERK pathways synergistically inhibited this response, suggesting co-ordinate regulation. Although from the present data alone we could not infer whether the Hsp27 pathway, which is an immediate pathway regulated by the up-stream kinase MK-2 regulated by MAPK^{p38}, is likely to exert a direct immunoregulatory effect on TNF-α biosynthesis, it is possible that Hsp27 might be implicated. This assumption is reinforced by recent evidence suggesting that Hsp27 can act as an endogenous protein circulating in the serum of breast cancer patients and a protein whose induction correlates with the onset of LPS shock (De *et al.*, 2000). Moreover, it has been reported that Hsp27 mediates regulatory effects on the biosynthesis of TNF-α through an IL-10 dependent mechanism, a pathway known to exert an anti-inflammatory role in several systems (Widmann *et al.*, 1999; De *et al.*, 2000).

Redox and ROS signalling regulating MAPK^{p38}-mediated induction of pro-inflammatory cytokines, and particularly TNF- α , has not been well characterized. It has been recently reported that ROS-initiated and redox-sensitive mechanisms converge on MAPK^{p38/ERK}-mediated regulation of TNF-a signalling in a cascade circuit, emphasizing the highly organized interactive nature of intracellular signals emanating from cell membranes leading to gene regulation and induction (De Keulenaer et al., 2000). Furthermore, a novel glutathione-sensitive antioxidant response element regulating the MAPK^{p38} cascade was reported in mediating the inducible expression of phase II enzymes, such as rGSTA2, known to be responsible for the protective adaptive responses to electrophiles and ROS (Kang et al., 2000). In a much more recent novel investigation, Hashimoto et al. (2001) reported a dual role for NAC in attenuating TNF-α-induced MAPK^{p38} activation and MAPK^{p38}-mediating IL-8 biosynthesis in vitro. In the present study, LPS-mediated secretion of TNF- α was attenuated by selectively blockading the MAPK^{p38/ERK}/MK-2 pathways and that MAPK-mediated regulation of TNF-a biosynthesis was shown to be ROS/redox-sensitive and modulated by the antioxidant and glutathione (GSH) precursor, NAC. NAC, a cysteine pro-drug (Aruoma et al., 1989; Bernard, 1991; Haddad et al., 2000), can suppress cytokine production (Jeannin et al., 1995; Matsumoto et al., 1998; Gosset et al., 1999; Haddad et al., 2001c) and protect against ROS-mediated lung injury (Bernard, 1991). The ratelimiting substrate for GSH biosynthesis is glutamate-cysteine $(K_m \text{ Glutamate} = 1.6 - 2 \text{ mM}; K_m \text{ Cysteine} = 0.3 \text{ mM})$ (Griffith & Meister, 1979; Meister, 1988; Aruoma et al., 1989; Bernard, 1991; Haddad et al., 2001c). Replenishing and sustaining intracellular GSH concentrations, therefore, is accomplished by administering compounds that increase the level of this amino acid (cysteine), or by promoting the activity of γ glutamylcysteine synthetase (γ -GCS), the rate-limiting enzyme in the biosynthesis of GSH (Meister, 1988). The ability of NAC to induce intracellular accumulation of GSH suggested that its potential to suppress TNF- α secretion residues on two possible pathways: Firstly, the amino acid cysteine provided by the administration of NAC eventually feeds into the biosynthetic machinery regulated by γ -GCS. GSH that is formed during this conversion detoxifies accumulating ROS through the glutathione-peroxidase coupled reaction (Haddad & Land, 2000a). Therefore, the pathway implicated with cysteine is to complement the biosynthetic process, where GSH can directly scavenge ROS. Secondly, NAC has antioxidant properties in that it is capable of directly scavenging and dissimulating accumulating ROS (Aruoma et al., 1989). We have previously shown that NAC induced intracellular formation of GSH (Haddad & Land, 2000a; Haddad et al., 2000), consistent with the observation reported in this study and that NAC ability to detoxify intracellular ROS blockaded the regulated downstream pathway for cytokine signalling (Haddad *et al.*, 20001a, c). Taken together, these data argue for ROS as potential messengers in regulating TNF- α signalling and that the ability of NAC to blockade this pathway resides in its potential either to deliver GSH, a major antioxidant thiol, or to detoxify excess ROS accumulation (Aruoma *et al.*, 1989; Bernard, 1991; Haddad & Land, 2000a).

This novel antioxidant potential of NAC was associated with the ability to suppress the phosphorylation and activation of the MAPK^{p38} pathway. Since ROS, derived from exposure to extracellular signals inducing stress, were reported to activate the MAPK^{p38} pathway (Su & Karin, 1996; Widmann et al., 1999), the likely possibility of NAC effectively acting as an antioxidant blockading this cascade is very probable, supported by the unequivocal observation that exogenous ROS up-regulated LPS-mediated MAPK^{p38} phosphorylation and selective antioxidants attenuated this effect. However, the possibility that NAC might be directly interacting with one or more of the components of the MAPK module and whether its blockading upstream kinases converging onto this pathway cannot be excluded. On the mechanism of action of NAC, as a thiol-modulating agent, in blockading the MAPK^{p38} pathway, we report the likely occurrence of several possible mechanisms. ROS, such as O_2^{-1} , H_2O_2 and OH may function as second messengers in signal transduction. Although MAPKs may be directly activated by oxidants, the role of ROS in the activation of the MAPKs pathway by cytokines is largely inferred on the basis of the inhibition of MAPKs by NAC and other antioxidants (Chan et al., 2001). It was also reported that NAC acts as an inhibitor of the c-Jun NH₂-terminal kinase (JNK) pathway regulated by TNF-a (Natoli et al., 1997) and that TNF- α stimulation of MAPK^{p42/44}, MAPK^{p38} and MAPK^{JNK} pathways was blockaded by NAC under serumfree conditions (Chan et al., 2001). The fact that the cellpermeable glutathione pro-drug, γ -GCE, was shown to be likewise potently effective in down-regulating LPS induced MAPK^{p38} phosphorylation, suggested that GSH is a major player in regulating this pathway and that NAC suppression of this response is GSH-sensitive. y-GCE is rapidly deesterified by intracellular esterase, thereby serving as an effective delivery agent for glutathione, which is a peptide incapable of crossing membranes in its native form. Although a distinction between the biological effects of γ -GCE and GSH is indiscriminate, intracellular conversion of y-GCE suggests that its effects are mediated by GSH. Exogenous/ endogenous glutathione, therefore, may feed into one of the well-characterized pathways of metabolism (Meister, 1988). For instance, GSH plays an important role in determining how readily pro-inflammatory genes can be regulated, and GSH/GSSG equilibrium is a major determinant of the activation of redox-sensitive transcription factors, including NF- κ B (Dröge *et al.*, 1994). Furthermore, apoptosis signalregulating kinase-1 (ASK-1) was recently identified as a MAP kinase kinase that activates the MAPK^{p38} pathway (Saitoh et al., 1998). Additionally, it was shown that TNF-a-mediated regulation of ASK-1 is ROS- and redox-sensitive and that NAC and thioredoxin (TRX), a redox molecule, blockaded the dimerization and the activity of ASK-1 (Saitoh et al., 1998; Hashimoto et al., 2001). Subsequently, it was

hypothesized that ASK-mediated regulation of the MAPK cascades depends on the interaction of ASK with the reduced form of a thiol modulating agent (NAC, GSH, TRX), thereby causing direct inhibition of the activity of the downstream pathway (Saitoh *et al.*, 1998). Although the activity of ASK-1 has not been assessed in this investigation, it is very likely that MAPK-mediated regulation of TNF- α biosynthesis in the alveolar epithelium involves a ROS/redox-sensitive mechanism regulating the activity of ASK-1 and probably other components of the converging up-stream cascades, thereby attenuating the stimulatory effect of MAPK pathway in TNF- α signalling. Collectively, the MAPK pathway seems to be directly involved in cytokine signalling and that MAPK-mediated regulation of LPS-induced TNF- α biosynthesis is ROS/redox-sensitive.

In summary, the results of the present investigation could be highlighted as follows: (i) Exposure to *E. coli*-derived LPS induced a time- and dose-dependent activation of the MAPK^{p38} pathway; (ii) LPS mediated regulation of MAPK^{p38} cascade was associated with the activation of the MAPKAP-K2 (MK-2) pathway, thereby allowing phosphorylation of Hsp27; (iii) Selective blockade of either MAPK^{p38} and MAPK^{ERK} pathway attenuated TNF- α biosynthesis induced by LPS, an effect synergistically amplified by simultaneous inhibition; (iv) Selective inhibition of NF- κ B attenuated, but

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did not abrogate, LPS-mediated secretion of TNF- α ; (v) LPS induced intracellular accumulation of ROS, which upregulated MAPK^{p38} phosphorylation, an effect attenuated by NAC and other antioxidants, including the GSH precursor, γ -GCE; (vi) NAC reduced LPS-induced TNF- α secretion, an effect associated with its ability to induce intracellular accumulation of GSH and lower than that of GSSG; and (vii) The antioxidant potential of NAC and its ability to feed cysteine onto the GSH biosynthetic machinery are likely to reside in its effectiveness to blockade the $MAPK^{_{p38}}$ pathway, thereby suppressing the downstream TNF- α signalling pathway. It is concluded that MAPKmediated regulation of LPS induced TNF- α biosynthesis in the alveolar epithelium is ROS and redox-sensitive and requires the involvement of GSH-mediated signalling pathways.

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