

Enhancement of neutrophil function by the bronchial epithelium stimulated by epidermal growth factor

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ABSTRACT: The bronchial epithelium is an important physical barrier that regulates physiological processes including leukocyte trafficking. The aim of the present study was to elucidate the mechanisms whereby the bronchial epithelium, stimulated by epidermal growth factor (EGF) as part of a response to acute or chronic injury, could activate and chemoattract human neutrophils.

Subconfluent human bronchial epithelial (16HBE) cells were stimulated with EGF to mimic the *in vivo* events after injury. The effect of the resulting EGF-conditioned media (CM) was compared with that of basal-CM with respect to neutrophil activation and chemotaxis. Such findings were then confirmed using primary bronchial epithelial cells (PBECs) from healthy volunteers.

EGF-CM from 16HBE cells caused increased expression of CD11b/CD66b and CD62L loss on neutrophils when compared with basal-CM. EGF-CM contained significant neutrophil chemotactic activity involving granulocyte-macrophage colony-stimulating factor and interleukin-8 that was potentiated by leukotriene B₄. This was dependent on neutrophil phosphatidylinositol-3-kinase activation and Akt phosphorylation, with partial regulation by phospholipase D, but not mammalian target of rapamycin. Consistent with these observations, EGF-CM derived from PBECs displayed increased chemotactic activity.

The present results suggest that the enhanced chemotactic activity of the epidermal growth factor-conditioned epithelium can enhance neutrophil-mediated immunity during acute injury, while during continued injury and repair, as in chronic asthma, this could contribute to persistent neutrophilic inflammation.

KEYWORDS: Bronchial epithelium, chemotaxis, epidermal growth factor, inflammation, lung, neutrophils

eutrophil-mediated inflammation is the subject of extensive research [1]. Influx of neutrophils into tissues begins by rolling, which is mediated via surface CD62L (L-selectin) interacting with complementary ligands on endothelial cells, and is followed by firm neutrophil adhesion to the endothelium using β_2 -integrins (CD11a and CD11b) and transmigration. Upon activation, neutrophils shed CD62L and increase surface expression of CD11b [2, 3] and CD66b [4]. This is accompanied by degranulation, respiratory burst [5, 6] and release of a range of proinflammatory agents [1, 5, 7]. Neutrophil-mediated inflammation occurs during acute infection, in acute respiratory distress syndrome [8] and in chronic diseases, such as chronic obstructive pulmonary disease [9]. Increasingly, neutrophils are being implicated in asthma [10], a disorder long viewed as an eosinophilic disease [11]. Thus, emerging evidence points to their role during infectious exacerbations and in severe chronic asthma [12–14]. Raised numbers are observed in bronchoalveolar lavage (BAL), induced sputum and both bronchial and transbronchial biopsies [7, 12, 15] with increased concentrations of neutrophilactive mediators interleukin (IL)-8 [13, 16], leukotriene (LT)B₄ [16, 17], granulocyte-macrophage colony-stimulating factor (GM-CSF) [18, 19] and tumour necrosis factor (TNF)- α [18, 20] in BAL.

The mechanisms regulating the accumulation and activation of neutrophils in the airways

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remain poorly understood but are increasingly thought to involve the bronchial epithelium and growth factors such as epidermal growth factor (EGF), which acts via the EGF receptor (EGFR), a member of the ErbB family (c-erbB1, HER1; c-erbB2, HER2, neu; c-erbB3, HER3; and c-erbB4, HER4) [21, 22]. In addition to its role in epithelial regeneration and repair [22], the EGFR pathway is believed to contribute to lung inflammation [23, 24]. Extensive immunostaining for EGFR in the asthmatic bronchial epithelium has previously been reported to be increased with disease severity and to correlate with IL-8 expression [25]. These data are consistent with observations that activation of the EGFR pathway causes IL-8 secretion [24, 26, 27] and provides a mechanism for neutrophil migration. EGFR-mediated mechanisms have also been implicated in bronchial epithelial repair in asthma [22, 28], augmenting the inflammatory potential of virusinfected bronchial epithelial cells [29] and modulating the effects of cigarette smoke [26]. Previous studies have reported that primary bronchial epithelial cell (PBEC) cultures derived from asthmatic patients release neutrophil-active mediators IL-8, GM-CSF and TNF-α [24, 30, 31]. Furthermore, there is evidence that EGF induces a corticosteroidinsensitive increase in IL-8 gene and protein expression by bronchial epithelial cells [24, 25] and a strong correlation exists between epithelial EGFR expression and both epithelial IL-8 immunostaining and neutrophil numbers has been demonstrated [24].

The functional significance of the augmented production of pro-inflammatory mediators and their relative roles in driving neutrophilic inflammation remain uncharacterised. Furthermore, little is known about the intracellular neutrophil signalling pathway(s) involved in the communication between the epithelium and neutrophils. The current authors hypothesised that exposure of bronchial epithelial cells to EGF induces pro-inflammatory factors that promote activation and recruitment of neutrophils and that this activation involves phosphatidylinositol-3-kinase (PI3K) and its downstream pathway. In healthy subjects, EGFRs are localised on the basolateral surfaces of epithelial cells where they are inaccessible to ligands in the airway lining fluid. However, damage to the barrier allows ligand penetration and subsequent EGFR activation. To test this hypothesis, the authors first used the human bronchial epithelial (16HBE) 14o-HBE cell line stimulated by EGF to mimic the activation that occurs when the epithelial barrier is damaged. Studies have shown that 16HBE cells display morphological, functional and permeability properties resembling those of the primary bronchial epithelium [32-34]. It was examined whether when the epithelium was switched into repair mode by EGF it could modulate neutrophil function by studying the ability of conditioned media from EGF-stimulated epithelium (EGF-CM) to activate and chemoattract neutrophils and to trigger signalling pathways in neutrophils. After showing major effects of EGFR on the way 16HBE cells modulate neutrophil function, the chemotaxis experiments were repeated using PBECs from healthy nonatopic individuals as a source of conditioned media (CM) to confirm that the observations seen in the cell line were reflective of the function of primary cells derived from volunteer subjects.

MATERIALS AND METHODS

Epithelial cell culture

The 16HBE epithelial cells and fresh PBECs, obtained by bronchial brushings during fibreoptic bronchoscopy from healthy nonatopic control subjects, were grown as previously reported [22, 35]. The 16HBE cells were cultured in T75 flasks in minimal essential media with Earle's salts (Gibco BRL, Paisley, UK) supplemented with 10% (v/v) foetal calf serum (FCS; TCS Biologicals, Buckingham, UK), 2 mM L-glutamine (Gibco BRL), 50 U·mL⁻¹ penicillin and 50 μg·mL⁻¹ streptomycin (Gibco BRL), whereas primary PBECs were grown in hormonally supplemented bronchial epithelial growth media (Clonetics, San Diego, CA, USA) with the primary passage 0 containing 0.5 ug·mL⁻¹ fungizone (Gibco BRL). Cell cultures were incubated at 37°C in a humidified atmosphere of 5% (v/v) CO₂ in air. Following sequential passages of epithelial cells (passage 2 for PBECs), $\sim 1 \times 10^5$ cells were seeded into 24-well culture plates (coated with 30 µg·mL⁻¹ type 1 collagen (PureCol, Inamed, Fremont, CA, USA) for PBECs) and grown to 70% confluency. The subconfluent cell monolayers were then rendered quiescent for 24 h in their relevant serum-free media (SFM); 16HBE cells were serum starved by culturing them with RPMI 1640 without phenol red containing 1% (v/v) bovine serum albumin (BSA), 2 mM L-glutamine, 50 U·mL⁻¹ penicillin and 50 μg·mL⁻¹ streptomycin, whereas PBECs were serum starved in bronchial epithelial basal media containing 1% insulin, transferrin and sodium selenite media supplement plus 0.3% (v/v) BSA for PBECs). To mimic the repairing and resting phenotypes, subconfluent epithelial cell cultures were incubated for 24 h with EGF (10 ng·mL⁻¹, 24 h) or SFM, respectively, and generated EGF-CM and basal-CM were harvested by centrifugation. Secretion of transforming growth factor- β_2 was increased significantly by EGF (see figure D1 in supplementary data) from 16HBE cells, consistent with the notion that EGF-induced repair causes release of epithelium-derived pro-fibrogenic factors [22]. The 16HBE epithelial cell CM were also analysed for GM-CSF, IL-8, LTB₄ and TNF-α levels using commercial ELISA kits (R&D Systems, Abingdon, UK).

Analysis of neutrophil activation induced by the EGFstimulated epithelium

Peripheral venous blood was obtained from normal healthy donors and neutrophils were purified as previously detailed [36, 37] using dextran sedimentation and discontinuous plasma-Percoll® gradients (GE Healthcare Ltd, Buckinghamshire, UK) with minor modifications. After separation, neutrophils were washed with 50% platelet-poor plasma-PBS solution, followed by Dulbecco's PBS with or without Ca²⁺ and Mg²⁺. This procedure routinely yielded >96% pure unprimed neutrophils, with <0.5% mononuclear cell contamination and >98% viability, as assessed by trypan blue exclusion.

In order to assess the effects of the epithelium on the expression of cell surface markers of neutrophil activation, freshly purified neutrophils (5×10^5 cells per condition) were suspended in PBS with Ca²⁺ and Mg²⁺ and incubated with epithelial CM at 37° C for 1, 5, 15, 30, 60 and 120 min. PBS or $10~\mu$ M *N*-formyl-methionyl-leucyl-phenylalanine (fMLP) were used as negative and positive controls, respectively. Reactions were quenched by adding 1 mL of ice-cold PBS, followed by immersion of tubes in ice. Cells were then washed in PBS and



incubated for 15 min on ice with 10% autologous serum to block nonspecific binding sites. Specific staining was then performed with fluorochrome-coupled anti-CD16, -CD14, -CD62L, -CD11b and -CD66b monoclonal antibodies (Ab; all BD Biosciences, Oxford, UK) using fluorescence-activated cell sorting (FACS) staining buffer (FSB; 2% (v/v) heat-inactivated FCS, 0.09% sodium azide in PBS) for 30 min on ice. Unbound Ab was removed by washing twice in FSB and cells were analysed on a FACS Calibur using Cell Quest software (both BD Biosciences). Neutrophils were identified by their forward and side-scatter characteristics and all gated cells were CD16+/CD14-. Dead cells were excluded by 7-amino-actinomycin D, the staining of which was routinely <5%.

To assess activation further, neutrophils were stimulated with CM for 30 min, the optimal time-point based on the timecourse study using flow cytometry, and assessed for secretion of myeloperoxidase (MPO) determined by the 3,3-dimethoxybenzidine method [37, 38]. Cells (10⁶) were first suspended in PBS and then incubated at 37°C with SFM, EGF, basal-CM or EGF-CM. Reactions were terminated by placing the plates on ice for 5 min before centrifugation ($200 \times g$ for 10 min at 4° C). Supernatants were incubated with phosphate buffer (pH 6.2), 3,3'-dimethoxybenzidine (0.2 mg·mL⁻¹) and H₂O₂ (90 nM) for 30 min at 37°C and reactions were terminated by adding 2 mM sodium azide prior to spectrophotometric detection of coloured products at an absorbance of 460 nm (Jenway 6505 UV/VIS spectrophotometer; Jenway, Felsted, UK). The amount of MPO released was expressed as percentage of the maximal MPO secretion obtained after stimulation with fMLP (10 μM).

Assessment of neutrophil chemotactic activity generated in bronchial epithelial culture

The ability of EGF-stimulated epithelial cells to generate chemoattractants for neutrophils was first assessed using 16HBE and then confirmed in PBECs from healthy volunteers. The chemotactic activity was measured in a fluorescence-based chemotaxis-chamber assay using calcein-labelled neutrophils as responder cells. Purified neutrophils were resuspended in fresh chemotaxis loading buffer (Hanks' balanced salt solution containing 10 mM HEPES (pH 7.4) and supplemented with 10% autologous serum) at a concentration of 10×10^6 cells·mL⁻¹, then incubated for 45 min at $37^{\circ}C$ with $5\,\mu M$ Calcein AM (Molecular Probes Europe BV, Leiden, the Netherlands). Either 31 µL of basal-CM, EGF-CM or SFM were loaded into the bottom wells of a 96-well plate chemotaxis chamber (ChemoTX® System; Neuroprobe, Gaithersburg, MD, USA). For inhibition studies, neutrophils were pretreated with the putative inhibitors (all from Calbiochem, Merck Biosciences Ltd, Nottingham, UK; CP-105696 was a gift from M. Yeadon, Pfizer Ltd, Kent, UK) for 30 mins prior to the addition to the upper wells. Subsequently, a polycarbonate filter (3-µm pore size) was placed over the bottom chamber. Neutrophils were washed with chemotaxis buffer to eliminate nonincorporated Calcein AM and loaded (30 µL, 3×10^5 cells·mL⁻¹) on top of the filter directly over each sample well in triplicate. After allowing cells to migrate at 37°C (5% (v/ v) CO₂) for 60 min, the nonmigrated cells were gently aspirated and 40 μL of detachment buffer (PBS supplemented with 20 mM EDTA and 0.3% (v/v) BSA) was applied on top of each well and incubated for 30 min at 4°C. Residual neutrophils that remained on top of the filter were flushed carefully with PBS

and the plate was centrifuged for 8 min at $120 \times g$ to collect any cells that had partially migrated through the filter. The ChemoTX® System was disassembled gently and the bottom plates were read in an FLX 800 Microplate Fluorescence reader (Bio-Tek, Winooski, VT, USA) using 485 nm and 530 nm as the excitation and emission wavelengths, respectively. Fluorescence of migrated neutrophils was extrapolated from a standard constructed curve using known numbers of fluorescently labelled cells. The number of cells migrating in the absence of chemoattractant was subtracted to determine chemoattractant-mediated chemotaxis.

Analysis of signalling via Akt by Western blotting

Neutrophils were exposed to basal-CM and EGF-CM from 16HBE cultures and fMLP (10 μM), SFM or EGF as controls for 5 min. Whole-cell protein lysates were then prepared from the cell pellets by solubilising in lysis buffer (20 mM Tris-HCl, pH 7.8, 138 mM NaCl, 1 mM EDTA, 1 mM sodium orthovanadate, 1% Nonidet P-40 and 1% Triton X-100) supplemented with a protease inhibitor cocktail (all reagents purchased from Sigma Aldrich, Poole, UK). Protein samples (50 µg of protein per condition) were denatured, separated by SDS-PAGE and transferred overnight onto nitrocellulose membrane by using the Mini-PROTEAN 3 apparatus (Bio-Rad Ltd, Watford, UK). Following the blocking step using PBS with 5% skimmed milk and 0.1% Tween 20 (Sigma Aldrich) for 1 h, the membranes were incubated with primary Ab at room temperature for 1 h with anti-phospho-Akt (Ser⁴⁷³ specific) monoclonal Ab (1:1,000 dilution; Cell Signaling Technology, Danvers, MA, USA). After sequential washing, the membranes were incubated for 1 h with the secondary Ab, a goat anti-mouse immunoglobulin G conjugated to horseradish peroxidase (1:2,500 dilution of stock, obtained from Sigma Aldrich). The immunoblotted membranes were visualised using Pierce ECL Western Blotting Substrate (Pierce, Cheshire, UK). For re-probing, membranes were stripped (in buffer containing 62.5 mM Tris, 2% SDS and 0.8% (v/v) β -mercaptoethanol) for 45 min at 55°C, followed by extensive washing before reblocking and re-probing with either anti-total-Akt polyclonal Ab (1:1,000 dilution; Cell Signaling Technology) or anti-α-tubulin (1:5,000 dilution; Sigma Aldrich).

Statistical analysis

All values are expressed as mean \pm SEM from a number of separate experiments. Data were analysed using an unpaired t-test. A p-value <0.05 was considered statistically significant.

RESULTS

Effects of epithelial culture-CM on neutrophil activation

Control conditions, *i.e.* SFM and basal-CM or EGF alone (data not shown), did not significantly alter the proportion of neutrophils expressing CD11b and CD66b until 2 h (fig. 1a and b). Incubation of neutrophils in EGF-CM markedly upregulated CD11b and CD66b at 30 min (p<0.001; fig. 1a and b). In parallel, CD62L expression markedly reduced within 5 min of exposure to both basal-CM and EGF-CM (fig. 1c) and followed a similar trend over 2 h. While the rate of CD62L shedding was similar for basal-CM and EGF-CM, this was significantly different throughout the time-course when compared with control conditions. EGF-CM also induced a significant increase in MPO release (p=0.0065). A further minor stimulatory effect on MPO release was observed when

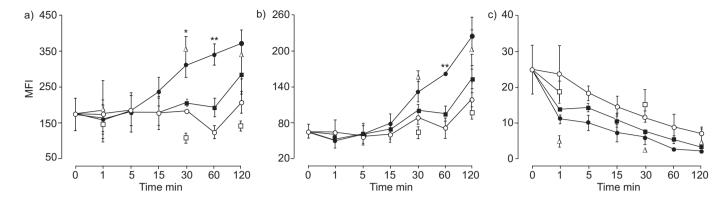


FIGURE 1. Time-course study of epidermal growth factor (EGF)-conditioned media (CM) effects on neutrophil activation marker expression in a) CD11b, b) CD66b and c) CD62L. Purified neutrophils were treated with serum-free medium (○), basal-CM (■) or EGF-CM (●). PBS (□) and *N*-formyl-methionyl-leucyl-phenylalanine (10 μM; Δ) were used as negative and positive controls, respectively. Expression of neutrophil surface molecules was determined by flow cytometric analyses, as described in the Materials and methods section. Data are presented as mean ± sem of four independent experiments performed using neutrophils purified from blood of four different donors. MFI: mean fluorescence intensity. *: p<0.05; **: p<0.01 compared with basal-CM-treated neutrophils.

EGF-CM-treated neutrophils were compared with basal-CM-treated cells, but this difference was not statistically significant (see figure D2 in the supplementary data).

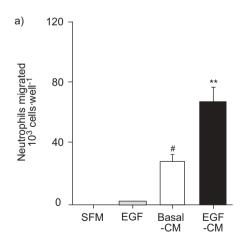
Neutrophil chemotactic activity detected in epithelial culture-CM

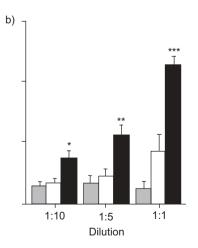
Both basal-CM and EGF-CM from 16HBE cells induced significant (p<0.0001) neutrophil migration (fig. 2a), with the effect of EGF-CM being significantly (p=0.0043) greater than basal-CM, while EGF alone had no effect (fig. 2a). Whereas dose–response curves of chemotactic responses to individual chemoattractants are typically bell-shaped, assessment of responses to three dilutions of EGF-CM (1:10, 1:5 and 1:1) showed that a 1:1 dilution of the original EGF-CM induced a

maximum migratory response (fig. 2b). Therefore, this dilution was chosen for all subsequent experiments. Using PBECs grown from bronchial brushings obtained from healthy volunteers, the effect of EGF on neutrophil chemotactic activity was shown to be similar, although both basal- and EGF-CM had a slightly less potent effect (fig. 2c).

Signalling pathways involved in the regulation of neutrophil chemotaxis induced by an EGF-CM

Based on previous studies [39, 40], it was hypothesised that the observed enhanced chemotactic response to EGF-CM involved activation of PI3K and its downstream signalling pathways in migrating neutrophils. Pre-treatment of neutrophils with the PI3K inhibitor, wortmannin (100 nM, 30 min), prior to exposure to





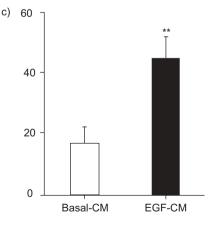


FIGURE 2. Chemotaxis induced by epidermal growth factor (EGF)-conditioned media (CM) and basal-CM. a) Calcein-loaded neutrophils were exposed to either serum-free medium (SFM; ■), EGF (■), basal-CM (□) or EGF-CM (■; all at 1:1 dilution). While EGF alone induced no chemotactic activity, EGF-CM contained greater activity than basal-CM. b) The epithelium-derived chemotactic activity acted in a concentration-dependent manner, with a maximal effect at 1:1 dilution. c) EGF-CM from human primary bronchial epithelial cells (PBECs) contained greater activity than basal-CM. Data are presented as mean ± sem of experiments performed in triplicate using neutrophils from five different donors and PBECs from bronchial brushings of six healthy control subjects. *: p<0.05; ***: p<0.01; ****: p<0.001 versus neutrophils treated with basal-CM alone; #: p<0.001 versus SFM-treated neutrophils.



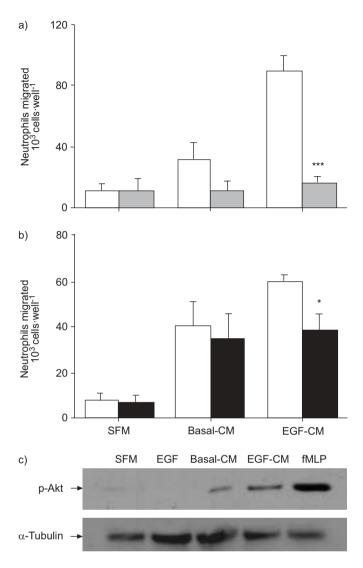


FIGURE 3. Effect of the a) phosphatidylinositol-3-kinase (PI3K) inhibitor wortmannin or b) the Akt inhibitor HIMO (1L-6-hydroxymethyl-chiro-inositol2-(R)-2-O-methyl-3-O-octadecylcarbonate) on neutrophil migration induced by epidermal growth factor (EGF)-conditioned media (CM) or basal-CM. Calcein-loaded neutrophils were pre-treated with either buffer (□), wortmannin (100 nM; ■) or HIMO (10 μM; ■) for 30 min before exposure of cells to serum-free medium (SFM), basal-CM or EGF-CM. Data are presented as mean ± SEM of seven and five independent experiments performed in triplicate, respectively. c) A Western blot of whole-neutrophil lysates showing that EGF-CM stimulation induces Akt phosphorylation. Purified neutrophils were stimulated with SFM, EGF, basal-CM, EGF-CM or N-formyl-methionyl-leucyl-phenylalanine (fMLP; 10 μ M) as a positive control at 37°C for 5 min, and then lysed. Equal amounts of protein from neutrophil lysates (50 µg of lysate per lane) were run on 10% SDS-PAGE, transferred to polyvinylidene fluoride and phosphorylated Akt (p-Akt) on Ser⁴⁷³ visualised by Western blotting. $\alpha\textsc{-Tubulin}$ expression is shown as a loading control. A representative blot of three independent experiments is presented. *: p<0.05; ***: p<0.001 versus untreated controls.

EGF-CM obtained with 16HBE cells reduced chemotaxis to levels achieved with basal-CM, with no effect on basal-CM-induced migration (fig. 3a). The role of Akt, a well characterised PI3K-dependent protein kinase known to regulate various neutrophil functions, was then investigated [41, 42]. While the selective Akt

inhibitor, 1L-6-hydroxymethyl-*chiro*-inositol2-(R)-2-O-methyl-3-O-octadecylcarbonate (10 μM , 30 min) [43], had no effect on basal-CM-mediated chemotaxis, it significantly reduced the EGF-CM-induced migration down to levels seen with basal-CM (fig. 3b). Western blotting using a monoclonal Ab specific for endogenously phosphorylated Akt (Ser^{473} specific) produced a distinct band of ${\sim}60$ kDa that corresponded to the predicted size of phosphorylated Akt (fig. 3c), with the intensity being greater in EGF-CM-stimulated cells when compared with basal-CM-treated neutrophils (fig. 3c).

Since phospholipase D (PLD) is essential for the hydrolysis of phosphatidylcholine into phosphatidic acid (PtdOH) and subsequent participation in neutrophil function [37], it was further hypothesised that the PLD enzymatic pathway may also regulate neutrophil migration induced by EGF-conditioned epithelium. The primary alcohol, butan-1-ol (0.5% (v/v)), which participates in a transphosphatidylation reaction by diverting specific PLD-derived PtdOH into a more stable metabolite whilst being a poor substrate for subsequent PLD-driven signalling cascades, caused significant inhibition, while, the control alcohol, butan-2-ol (0.5% (v/v)) had no effect (fig. 4a).

Finally, the current authors examined whether EGF-CM-mediated neutrophil chemotaxis occurs *via* specific members of the PI3K-related kinase family, most notably mammalian target of rapamycin (mTOR), a target substrate of the PI3K/Akt signalling pathway [44]. Using rapamycin, which selectively inactivates mTOR, it was shown that chemotaxis induced by either EGF-CM or basal-CM was largely unaffected by a maximal inhibitory concentration (100 nM) of this lipophilic macrolide (fig. 4b).

Characterisation of neutrophil-active factors released by epithelial cells

Initial time-course analysis of CM from 16HBE cells showed that IL-8 release peaked at 24 h (fig. 5a). Analysis of the same time-point for the other mediators showed that, by comparison with basal-CM, EGF-conditioned epithelial cells also secreted greater quantities of GM-CSF (334.8 \pm 116.83 versus 85.72 \pm 13.4 $pg\cdot mL^{-1}$), TNF- α (54.17 \pm 11.07 versus 26.07 \pm 1.9 $pg\cdot mL^{-1}$; p<0.05) and IL-8 (341.24 \pm 48.49 versus 151.5 \pm 27.29 $pg\cdot mL^{-1}$; p<0.001; fig. 5b–d). In contrast, the EGF-conditioned epithelium did not produce any more LTB₄ than the control epithelium (40.29 \pm 3.81 and 33.11 \pm 4.9 $pg\cdot mL^{-1}$ in basal-CM and EGF-CM, respectively; fig. 5e).

The roles of the individual neutrophil chemotaxins were studied further using neutralising antibodies and blocking agents. In preliminary experiments, chemotactic dose–responses were obtained for TNF- α , GM-CSF, IL-8 and LTB₄ (see figure D3a–d in the supplementary data). In addition, the effects of neutralising monoclonal Abs against TNF- α and GM-CSF, and the selective antagonists for the IL-8 receptor, CXCR2 (SB-225002), and the LTB₄ receptor, BLT1 (CP-105696), were assessed (see figure D3e–h in the supplementary data). Using optimal concentrations of blocking agents, the chemotactic activity in basal-CM could not be significantly inhibited by any antagonist or neutralising monoclonal Ab in isolation. The activity was partially, but not significantly, reduced only when either the selective BLT1 receptor antagonist or the CXCR2 antagonist were combined with monoclonal Abs for TNF- α or GM-CSF

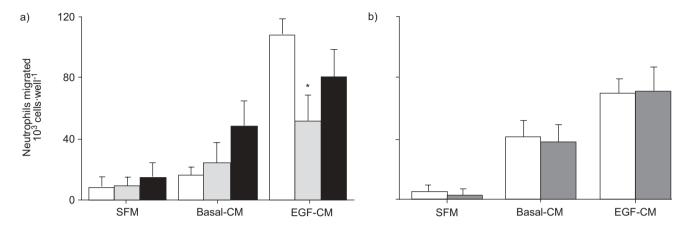


FIGURE 4. Effect of the modulators of a) phospholipase D or b) the mammalian target of rapamycin inhibitor, rapamycin, on basal-conditioned media (CM) and epidermal growth factor (EGF)-CM-triggered neutrophil migration. Calcein-loaded neutrophils (10 × 10⁶ cells·mL⁻¹) were pre-treated with either buffer (□), butan-1-ol (0.5% (v/v); ■), butan-2-ol (0.5% (v/v); ■) or rapamycin (100 nM; ■) for 30 min before exposure of cells to serum-free medium (SFM), basal-CM or EGF-CM for 60 min at 37°C. The number of chemoattractant-mediated neutrophils was extrapolated as detailed in the Materials and methods section. Data are presented as mean±sem of four and six independent experiments performed in triplicate, respectively. *: p<0.05 versus untreated controls.

(fig. 6a). EGF-CM-mediated chemotaxis could be partially attenuated close to basal levels when the CXCR2 (45.2% inhibition compared with control neutrophils) and BLT1 receptor (51.2% inhibition) antagonists, or the anti-GM-CSF monoclonal Ab (56.3% inhibition) were applied individually. Anti-TNF- α monoclonal Ab alone was ineffective (fig. 6b). Complete abrogation of neutrophil chemotaxis caused by EGF-CM to below basal levels (present in basal-CM) was achieved by combining anti-GM-CSF monoclonal Ab with either the CXCR2 or BLT1 antagonists alone or together (fig. 6b). Interestingly, the co-application of anti-TNF- α monoclonal Ab to either anti-GM-CSF monoclonal Ab or the CXCR2 and BLT1 antagonists partially reversed some of the inhibitory effects achieved with these blocking agents alone (fig. 6b).

To confirm further that the chemotactic activity present in EGF-CM involves LTB₄, 16HBE cells were cultured in the presence of the 5-lipoxygenase-activating protein (FLAP) inhibitor, MK-886 (1 μ M), for 30 min prior to EGF stimulation. The increase in chemotaxis towards EGF-CM was significantly attenuated by MK-886 to levels seen with basal-CM (fig. 7a), consistent with its effect on blocking 5-lipoxygenase (5-LO)-mediated LTs, suggesting that LTs participated in the generation of the chemotactic activity.

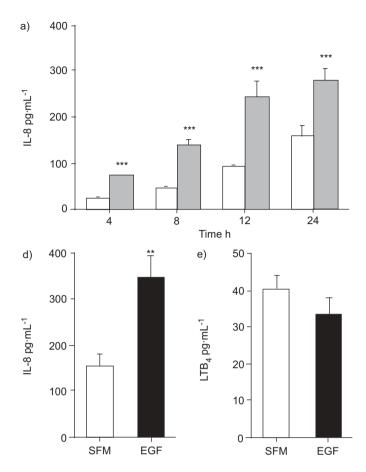
Having observed that upon pre-treatment of neutrophils with CP-105696 neutrophil migration to EGF-CM was substantially reduced, it was questioned whether this was due to the BLT1 antagonist blocking the chemotactic effects of LTB₄ generated by migrating neutrophils upon their initial exposure to EGF-CM. Indeed, 5-LO-initiated pathways and their products, including LTB₄, have been previously reported to be markedly increased in neutrophils during their migration in a murine air pouch-model of inflammation [45]. Therefore, the FLAP inhibitor, MK-886 (1 μM, 30 min), was used to analyse whether

neutrophil migration is mediated *via* autocrine release of LTB₄ upon exposure to basal-CM or EGF-CM. Neutrophil migration in response to both CM after pre-treatment with MK-886 remained unaltered (fig. 7b). MK-886 was also ineffective in diminishing neutrophil chemotaxis towards IL-8 and LTB₄ (fig. 7b), excluding the possibility of an autocrine process involving LTB₄ and suggesting that steady-state levels of LTB₄ can synergise with other critical mediators in EGF-CM.

DISCUSSION

Previous studies [24, 26, 27] have shown that, when stimulated by EGF, the epithelium produces mediators that activate neutrophils, but there has been no direct evidence to link these mediators with activation and chemotaxis of neutrophils and to quantify their relative contribution. The current study presents evidence that, while a resting epithelium can influence, to an extent, neutrophil behaviour, these responses are considerably enhanced when epithelial cells are stimulated by EGF. This setting reflects a response to any type of acute or chronic epithelial damage that compromises the physical integrity of the epithelial barrier allowing exposure of EGFRs on the basolateral surface of the epithelium to ligands present in the airway lining fluid. The present study has shown that neutrophil chemotactic responses to EGF-CM are regulated through PI3K and downstream signalling pathways involving Akt and PLD but not mTOR. It also highlights that the activated epithelium possesses a greater secretory potential as it releases larger quantities of the neutrophil-active mediators, GM-CSF, IL-8 and TNF-α. Consequently, the inhibition of their activity resulted in complete abrogation of the additional chemotactic activity that the EGF-conditioned epithelium produces over and above levels seen with a resting epithelium. Finally, the current authors have also demonstrated that while LTB₄ release by the activated epithelium is not raised relative





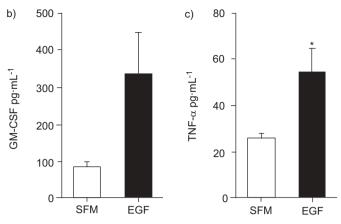


FIGURE 5. Treatment of human bronchial epithelial (16HBE) cells with epidermal growth factor (EGF) upregulates epithelium-derived mediator release. a) Time-course of interleukin (IL)-8 production measured by ELISA in conditioned media (CM) harvested at the indicated time-points after stimulation of 16HBE cells with serum-free medium (SFM; \square) or EGF (10 ng·mL; \blacksquare). The CM were also analysed for b) granulocyte-macrophage colony-stimulating factor (GM-CSF), c) tumour necrosis factor (TNF)- α , d) IL-8 and e) leukotriene (LT)B₄ secretion at the 24-h time-point. Apart from LTB₄, the levels of all mediators detected in EGF-treated cells (\blacksquare) were considerably higher than from SFM-treated cells (\square). Data are presented as mean \pm seM of four to nine separate experiments, each performed in duplicate. *:p<0.05, **: p<0.01, ***: p<0.001 versus SFM-treated controls.

to the basal epithelium, it acts to potentiate the chemotactic responses of the other pro-inflammatory mediators present in EGF-CM. Taken together, the results of this study show that the epithelium has a significant potential to enhance neutrophil function when driven into a repair mode, a phenotype that is seen after acute (e.g. viral infection) or chronic (e.g. asthma) lung injury.

The present study demonstrates for the first time that the activated epithelium can upregulate CD11b and CD66b, and reduce CD62L expression on neutrophils. Neutrophils utilise such processes in the acute setting when enhanced neutrophil function is beneficial at a time when the epithelial barrier is compromised. Following restoration of the epithelial barrier, EGF in the mucosal lining fluid is unable to stimulate basolaterally localised EGFRs in the pseudostratified columnar bronchial epithelium. If, however, the insult is excessive and/or chronic and does not allow complete repair, EGF-mediated epithelial activation will persist, causing sustained neutrophil activation with the potential to cause further tissue damage [5, 7, 10, 46]. A pivotal role of EGF in repair processes has been demonstrated in inflammatory bowel diseases [47, 48], renal injury [49], psoriasis [47, 50] and corneal damage [47, 51]. While these disorders also have a significant neutrophilic component, the association between EGF-mediated epithelial repair and neutrophil activation and recruitment has not been explored in detail. Hence, the demonstration of EGF modulatory effects in human PBECs herein complements and extends previous observations [24, 25], further implicating interaction between the repairing bronchial epithelium and neutrophils as a mechanism of neutrophil-mediated tissue damage.

In agreement with previous reports of neutrophil migration induced by various chemoattractants [39, 40], a role for PI3K signalling in the activation and migration of neutrophils induced by an EGF-conditioned epithelium was demonstrated. Likewise, EGF-CM-mediated neutrophil chemotaxis also involved enhanced activity of Akt downstream of PI3K and the PLD signalling pathway, with the latter finding being consistent with the report by LEHMAN et al. [52]. However, selective inhibition of mTOR did not affect the migratory responses to either basal-CM or EGF-CM, which is surprising given the reports by GOMEZ-CAMBRONERO [44] that neutrophil migration induced by GM-CSF is dependent on mTOR. This apparent discrepancy may arise because of regulation of neutrophil chemotaxis occurring via an mTOR component that is resistant to rapamycin treatment. This premise is based on observations in various mammalian cell types that mTOR can remain functionally active even after rapamycin treatment [53, 54]. This large protein kinase exists in two distinct complexes: one contains mTOR, GBL and raptor, and can be inhibited by rapamycin [55], while the other contains mTOR, GβL and rictor, and is rapamycin insensitive [53, 55]. The cellular function of the latter complex is only beginning to be unravelled [53] and the current study did not address the question of whether the rictor-mTOR complex regulates the

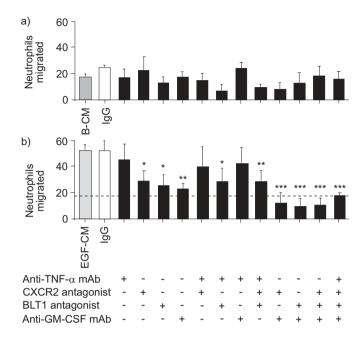


FIGURE 6. Neutralisation/blockade of epithelial-derived mediator activity abrogates neutrophil migration (10^3 cells·well-1). a) Administration of the BLT1 antagonist, CP-105696 ($10~\mu\text{M}$) or the CXCR2 antagonist, SB-225002 (100~nM), along with the anti-tumour necrosis factor (TNF)-α monoclonal antibody (mAb; $100~\mu\text{g·mL}^{-1}$) or anti-granulocyte-macrophage colony-stimulated factor (GM-CSF) mAb ($100~\mu\text{g·mL}^{-1}$), partially attenuated basal (B)-conditioned media (CM)-mediated chemotaxis compared with the untreated control neutrophils. b) Complete abrogation of the chemotactic activity exerted by epidermal growth factor (EGF)-CM on neutrophils was achieved down to basal levels (- - - - -) with the combination of anti-GM-CSF mAb with SB-225002 and/or CP-105696. Data are presented as mean ± SEM of five independent experiments performed in triplicate using peripheral blood neutrophils isolated from different donors. Ig: immunoglo-bulin. *: p<0.05; **: p<0.01; ***: p<0.001 versus untreated EGF-CM control.

observed rapamycin-insensitive chemotaxis in response to basal-CM and EGF-CM.

The present authors found that bronchial epithelial cells increase GM-CSF, IL-8 and TNF-α production when stimulated by EGF. Using selective inhibitors, it was possible to dissect the relative contributions of these epithelium-derived factors and to show synergy between individual mediators. It has been reported previously that neutrophilic chemotaxins, such as IL-8, are markedly upregulated in the altered epithelial phenotype in asthma [24]. A more recent study [27] has shown that the EGFR pathway is involved in stimulating IL-8 production in airway epithelial cells via a novel signalling cascade involving activation of TNF-α converting enzyme by dual oxidase 1. However, this chemokine is unlikely to be solely responsible for driving epithelial-mediated neutrophil migration as the present data clearly show that when the selective antagonist for the CXCR2 receptor (SB-225002) was applied individually, it failed to completely abrogate neutrophil chemotaxis evoked by EGF-CM (fig. 6b). Furthermore, EGF significantly upregulated secretion of GM-CSF and the observed chemotactic responses were sensitive to both the neutralising anti-GM-CSF monoclonal Ab and the selective BLT1 antagonist, suggesting that these two mediators have independent effects on the mediation of EGF-CM-driven chemotaxis.

Bronchial epithelial cells have been previously shown to generate biologically active LTB₄ [56, 57]. A recent study has shown that both 16HBE cells and PBECs constitutively express 5-LO, FLAP and LTA₄ hydrolase at the mRNA and protein levels [58]. As detected by ELISA and confirmed by reverse-phase high-performance liquid chromatography purification, the amounts of LTB₄ released over 6 h by stimulated 16HBE cells (119 pg per 10⁶ cells) were essentially similar to LTB₄ levels of 30 pg per 10⁵ cells detected at 24 h in the current

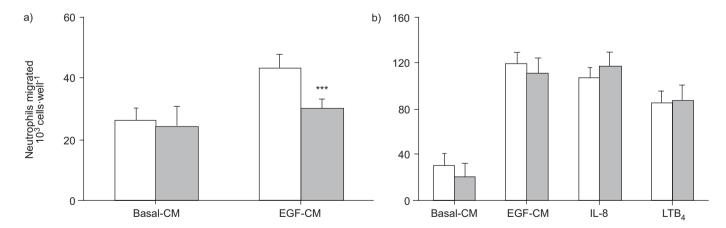


FIGURE 7. a) Inhibition of leukotriene synthesis in reparative human bronchial epithelial (16HBE) cells using the 5-lipoxygenase-activating protein (FLAP) inhibitor, MK-886, attenuates epidermal growth factor (EGF)-conditioned media (CM)-induced neutrophil migration. 16HBE cells were pre-treated with either buffer (□) or MK-886 (1 μM; ■) for 30 min before stimulation with serum-free medium or EGF (10 ng·mL⁻¹) for 24 h at 37°C. CM was collected from epithelial cultures then the chemotaxis was assessed in calcein-loaded neutrophils (10x10⁶ cells·mL⁻¹). b) Autocrine secretion of leukotriene (LT)B₄ does not regulate EGF-CM-mediated chemotaxis as inhibition of FLAP in neutrophils using MK-886 does not affect the migratory responses. Calcein-loaded neutrophils (10x10⁶ cells·mL⁻¹) were pre-treated with buffer (□) or MK-886 (1 μM; ■) for 30 min before exposure of cells to basal-CM, EGF-CM, interleukin (IL)-8 (10 nM), or LTB₄ (100 nM) for 60 min at 37°C. Data are presented as mean±sem of four independent experiments performed in triplicate using neutrophils purified from blood of different donors. ***: p<0.001 versus untreated controls.

study. Both recent data [58] and the current study show that LTB₄ can be significantly reduced by pre-treatment with the specific FLAP inhibitor MK-886, confirming the authenticity of the measured LTB₄. The current study further demonstrated bioactivity of the LTB4 released by 16HBE cells because selective blockade of the BLT1 receptor using CP-105696 significantly reduced EGF-CM-mediated chemotaxis and this LT was also able to potentiate the effects of the other chemotactic factors present in EGF-CM (fig. 6b). Of note, the subnanomolar concentrations of LTB4 present in EGF-CM (~30 pg·mL⁻¹, as detected by ELISA; fig. 5e) were at an adequate level to induce chemotaxis of human neutrophils [59]. Comparison of the results obtained with basal-CM and those with EGF-CM showed no differences in amounts of LTB₄ release, indicating that EGF is not capable of modifying the spontaneous release of LTB4 from human PBECs described previously, at least in 24-h cultures.

In conclusion, this is the first in-depth elucidation of direct modulatory effects of an epidermal growth factor-conditioned epithelium on human neutrophil activation and migration. The current authors speculate that while epidermal growth factor has the capacity to enhance neutrophilic innate immunogenicity during acute lung damage (e.g. viral injury), an aberration of these responses may be pathological in the context of chronic diseases (e.g. severe neutrophilic, forms of asthma) where bronchial epithelial damage is a common feature.

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