Neutrophil Elastase Causes MUC5AC Mucin Synthesis Via EGF Receptor, ERK and NF-κB Pathways in A549 Cells

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Background: Neutrophil elastase (NE) was found to increase the respiratory mucin gene, MUC5AC, although the molecular mechanisms of this process remain unknown. We attempted to determine the signal transduction pathway through which NE induces MUC5AC gene expression in bronchial epithelial cells.

Methods: A fragment of 1.3 Kb MUC5AC promoter which had been cloned into the pGL3-Basic luciferase vector was transfected to the A549 cells. By measuring the luciferase activity, we were able to evaluate the MUC5AC promoter activity in A549 cells. The involvement of mitogen-activated protein kinases (MAPK) was confirmed by Western blotting. To confirm the involvement of nuclear factor-kappaB (NF-κB), we used site-directed mutagenesis and electrophoretic mobility shift assay (EMSA) autoradiogram. The MUC5AC mRNA expression was confirmed by RT-PCR.

Results: NE increased the transcriptional activity of the MUC5AC promoter in A549 cells. The increased transcriptional activity of the MUC5AC promoter by NE was found to be associated with increased NF-κB activity. Site-directed mutagenesis showed that the transfection of the mutated NF-κB binding sites from the PGL3-MUC5AC-3752 promoter luciferase reporter plasmid decreased the luciferase activity after NE stimulation. Among the MAPKs, only extracellular signal-regulated kinases (ERK) were involved in this NE-induced MUC5AC mucin expression. RT–PCR also showed that NE increased MUC5AC mRNA. An EMSA autoradiogram revealed that NE induced NF-κB:DNA binding.

Conclusions: These results indicate that human NE induces MUC5AC mucin through the epidermal growth factor receptor (EGF-R), ERK, and NF-κB pathways in A549 cells.

Key Words: Leukocyte Elastase, Mucins, Extracellular Signal–Regulated MAP Kinases, NF-κappaB

INTRODUCTION

Goblet cell hyperplasia is a prominent feature in many chronic obstructive airway diseases associated with mucus hypersecretion, including chronic bronchitis, bronchiectasis, cystic fibrosis, and bronchial asthma. In these diseases of the airway, neutrophils are recruited into the airways. Hypersecretion from an increased number of goblet cells has been considered to contribute to mucus plugging and airway obstruction1. In the secretions of the airway, MUC5AC is a major mucin which is secreted from the goblet cells of the airway surface epithelium2.
Mucins are large molecular weight glycoproteins which are composed of 10 to 20% protein and 80 to 90% complex carbohydrates. Neutrophils release elastase, a serine protease that is found in high concentrations in the airway surface fluids of patients with chronic obstructive airway diseases. Purified neutrophil elastase (NE) was previously shown to be a potent secretagogue for goblet cells in vitro. In one report, NE increased MUC5AC mucin gene expression and glycoprotein production in respiratory epithelial cells: this increase in MUC5AC gene expression is due to the stability of MUC5AC mRNA stability. In contrast, the induction of the MUC5AC gene by NE is also related to the reactive oxygen species.

Mitogen-activated protein kinase (MAPK) involves NE–induced morphological changes in human bronchial epithelial cells. Mammals express at least four groups of MAPK: extracellular signal-related kinases (ERK)–1 and –2, p38 kinases, ERK–5, and c-Jun amino-terminal kinases (JNK 1, 2 and 3), which are also known as stress–activated protein kinases, p38 kinases and ERK–5. MAPKs are a family of proline–targeted, serine–threonine kinases that transduce environmental stimuli to the nucleus. In this report, we sought to determine the signaling pathway of the human neutrophil elastase (HNE)–induced MUC5AC gene expression in the airway epithelial cells of the airway. We demonstrated that ERK is involved in elastase–induced MUC5AC mucin expression, and also found that epidermal growth factor receptor (EGF–R) and nuclear factor kappaB (NF–κB) are also involved in this signal transduction pathway.

MATERIALS AND METHODS

Cell culture and stimulation

A549 cells, a human pulmonary adenocarcinoma cell line, were grown on 100 mm dishes (Nunc, Denmark) in RPMI 1640 medium containing 10% fetal bovine serum (FBS), penicillin (100 U/mL), streptomycin (100 μg/mL), and HEPES (25 mM) at 37°C in a humidified, 5% CO₂, water-jacketed incubator. At confluence, the A549 cells were maintained for 48 hours in 6–well culture plates at 2×10⁵ cells/well. The cells were then starved of serum for an additional 24 hours and stimulated with NE. MUC5AC promoter activity was determined by measuring luciferase after the lysing of the transfected cells and normaliztion by co–transfection with the β-galactosidase expression plasmid, pβgal-control vector (Clontech). β-galactosidase activity was measured in the luminometer (Turner Designs) in accordance with the instructions of the manufacturer of the pβgal-control vector (Clontech). All transfections were performed in triplicate wells: results were reported as emitted light per well (mean±S.D).

RT–PCR

RNA was isolated from cell cultures as previously described, using an RNA isolation kit (TRIZOL reagent, Invitrogen). The RNA was then reverse transcribed into cDNA. Oligonucleotide primers were designed on the basis of published sequences for human MUC5AC. The primers for amplifying MUC5AC cDNA were 5’–TCCGGCTCTACCTTCTCC–3’ (5’–primer) and 5’–ACTTGGGCACCTGGTGCTG–3’ (3’–primer). The primers for amplifying glyceraldehyde–3–phosphate dehydrogenase (GAPDH) were 5’–ACCAAGTCCATGCACCCAC–3’ (5’–primer) and 5’–TCACACCCGTGTTGCTGTA–3’ (3’–primer). PCR was performed using the standard technique. The amplification efficiency for MUC5AC was compared with GAPDH, and their ratio was calculated. Statistical comparisons were made using Student’s t test.
Table 1. Sequences of the pairs of oligonucleotides used in PCR to produce deletion mutants of the NF-κB binding sites in the MUC5AC gene

| Deletion mutant | Orientation | Oligonucleotide used for site-directed mutagenesis (5' -> 3') |
|-----------------|-------------|------------------------------------------------------------|
| NF-κB : Mutant 1| MS          | CTATCTGTGCTGGGACTGCTGCTGCTGCAAGTCAA                      |
|                 | MAS         | TTCTGTGCTGGGACTGCTGCTGCTGCAAGTCAA                        |
|                 | Normal      | GCCCTCTGTGCTGGGACTGCTGCTGCTGCAAGTCAA                    |
| NF-κB : Mutant 2| MS          | GACCTTTCTGTGCTGGGACTGCTGCTGCAAGTCAA                    |
|                 | MAS         | GACCTTTCTGTGCTGGGACTGCTGCTGCAAGTCAA                    |
|                 | Normal      | GACCTTTCTGTGCTGGGACTGCTGCTGCAAGTCAA                    |
| NF-κB : Mutant 3| MS          | GTATGTGCGGAGGAGCCCGCTGAGGCAGGCAAGTCAA               |
|                 | MAS         | GTATGTGCGGAGGAGCCACCTGAGGCAGGCAAGGCT               |
|                 | Normal      | GTATGTGCGGAGGAGCCCGCTGAGGCAGGCAAGGCT               |
| NF-κB : Mutant 4| MS          | GGGGATGATGTGCTGGGAGCCGGAGGCAAGGCAAGGCT |
|                 | MAS         | GGGGATGATGTGCTGGGAGCCGGAGGCAAGGCAAGGCT |

MS, mutant sense; MAS, mutant antisense.

Immunoctochemistry

Cells grown for 48 hours on 6-chamber slides were stimulated with NE (50 nM) and fixed with 100% methanol for 5 minutes. Cells were incubated with mouse monoclonal antibody (Ab) to MUC5AC (clone 45M1, 1:1,000, New Markers, Fremont, CA) for one hour and were then washed four times with phosphate buffered saline (PBS) to remove excess primary antibody. After one hour, plates were washed three times with PBS. The plates were then incubated with biotinylated horse anti-mouse IgG at a 1:2,000 dilution for 30 minutes. Bound antibodies were visualized according to standard protocols for the avidin–biotin–alkaline phosphatase complex method.

Site-directed mutagenesis

The MUC5AC gene fragments for the transcription factor NF-κB binding sites were mutated in order to determine which part is necessary for the initiation process of MUC5AC gene transcription. Two known portions of NF-κB binding sites (base pairs −973 to −939 and −237 to −203) were present in the MUC5AC promoter. Human MUC5AC regulatory regions were mutated (M1: −971 to −939, M2: −973 to −941, M3: −235 to −203, M4: −237 to −205) using a mutagenesis kit (Quick-Change Site-directed Mutagenesis Kit; Stratagene) (Table 1). PCR was performed by incubating 1 μL of pGL3 MUC5AC-3752 promoter luciferase reporter plasmid (10 ng/L) with 130 pmole of upstream and downstream primer, 400 μM of total dNTP, 3 μL of Quick solution, and 2.5 units of DNA polymerase in a final volume of 50 μL. PCR was carried out for 10 cycles (30 seconds at 95°C, 60 seconds at 55°C, 60 seconds at 72°C) and finished with an extension at 72°C for 10 minutes. After PCR, 1 μL of restriction enzyme, Dpn I (10 U/μL), was added to each sample, which was then incubated at 3 7°C for one hour. With each mutant, cell transformation to the E.coli was performed with an XL10-Gold E. coli kit (Stratagene). After sequencing, mutated DNA was harvested with Quiagen Midi-prep.

Western blot

A549 cells grown in 6-well culture plates were starved of serum for 24 hours, and were then stimulated with HNE for 30 minutes. In inhibition studies, a selective EGF-R tyrosine kinase inhibitor (AG1478: 50 μM), an MEK inhibitor (PD98059; 50 μM), and an NF-κB inhibitor (CAPE: 20 μg/mL) were added to the medium 30 minutes prior to stimulation with HNE. Cells were then lysed on ice in PBS lysis buffer containing 50 mM Tris–HCl, pH 7.5, 150 mM NaCl, 1% Nonidet P-40, 0.5% sodium deoxycholate, 0.1% SDS, 1 mM EDTA, 0.2% sodium azide, 1 μg/mL aprotinin, 1 g/mL pepstatin A, 0.2 mM phenylmethylsulfonyl fluoride (PMSF), and 0.6 mM sodium vanadate. The lysate proteins were mixed with the same amount of loading dye, and were then denatured with heating at 95°C for 10 minutes. The sample proteins (20 μL) were separated by SDS–PAGE in 10% acrylamide gel. The resulting gel was equilibrated in the transfer buffer (48 mM Tris-base, 39 mM glycine, 20% methanol, 0.037% SDS, pH 8.3). The proteins were then electrophoretically transferred to polyvinylidene difluoride (PVDF) membranes (Bio-Rad Laboratories), after which the membranes were blocked for one hour with 3% skim milk containing 20 mM Tris–HCl, 150 mM NaCl, and 0.05% Tween 20, at pH 7.5 at room temperature. The membranes were then incubated overnight with primary antibody (1:1,000) in 2% skim milk/Tris–buffered saline with Tween (TBST) buffer. The membranes were washed three times with TBST solution and incubated with horseradish peroxidase–conjugated rabbit
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Figure 1. Neutrophil elastase (NE) increases the transcriptional activity of MUC5AC promoter. MUC5AC reporter plasmid was transfected to the A549 cells, which then underwent treatment with vehicle or NE for 30 minutes at concentrations of 10, 25, 50, 100, and 200 nM concentrations. The cells were then harvested for measurement of luciferase activities. Significant difference in the transcriptional activity of MUC5AC promoter between control and 50 nM of NE stimulation to transfected A549 cells was detected are found. Values represent the average of six replicates normalized to co-transfected \( \beta \)-galactosidase luciferase activity, with standard deviation indicated with error bars (*, \( p < 0.05 \) compared with transfection with MUC5AC promoter without NE treatment).

Figure 2. Effects of inhibitors on the MUC5AC promoter activity in A549 cells. The nuclear factorkB (NF-kB) inhibitor (CAPE; 20 \( \mu \)g/mL), MAPK/ERK kinase (MEK) inhibitor (PD98059; 50 \( \mu \)M), and EGF-R inhibitor (AG1478; 50 \( \mu \)M), were added before treatments with neutrophil elastase (NE) (50 nM); all of these inhibitors markedly inhibited the promoter activity of MUC5AC in the A549 cells. Values represent the average of four assays, and standard deviation is provided by error bars (*, \( p < 0.01 \) compared with transfection with MUC5AC promoter without NE treatment).

Electrophoretic mobility shift assay (EMSA)

Transfected A549 cells were stimulated with HNE (50 nM) for 30 minutes. The cells were washed with ice-cold PBS and centrifuged at 3,000 rpm for five minutes. Buffer A (0.9 M HEPES [pH 7.9], 1 M MgCl\(_2\), 2 M KCl, 100 mM DTT, 100 mM PMSF) was added to the cell pellets: the cell pellets were kept on ice for 10 minutes. The cell pellets were again purified after centrifugation at 3,500 rpm and 12,000 rpm, each for five minutes. Buffer C (0.5 M HEPES [pH 7.9], 1 M MgCl\(_2\), 2.5 M NaCl, 100 mM EDTA, 100 mM DTT, 4% glycerol) was added to the cell pellets: the cell pellets were allowed to react on ice for 20 minutes. Nuclear extracts were finally isolated after centrifugation at 12,000 rpm for 10 minutes. The primers for EMSA were constructed (sense: 5'-TGTGCTGGGACTGCTCGGA CCA-3'; antisense: ACACGACCCTGACGAGCCTGGT) and heated at 95°C for 10 minutes after mixing the upstream and downstream primers at 1 nmole. The oligonucleotide probes were end-labeled with \( \gamma \)-p\(^{32}\)dATP (3,000 Ci/mmole at 10 mCi/mL: Amersham Co., Arlington Heights, IL) in the presence of T4 polynucleotide kinase and were isolated after being passed through a Sephadex G-25 column. Binding reactions containing equal amounts of nuclear protein extract (10 \( \mu \)g) and 35 fmol (~ 50,000 cpm, Cherenkov counting) of radio-

**RESULTS**

Neutrophil elastase increases the transcriptional activity of MUC5AC promoter

Before MUC5AC promoter transfection, transcriptional activity was nearly undetectable in a luminometer assay. After transfection, luciferase activity was found to have increased, and
Figure 3. Characterization of the nuclear factor-kB (NF-κB) site and NF-κB in MUC5AC mucin induction by neutrophil elastase (NE). Two portions of the NF-κB binding sites of MUC5AC reporter plasmid were mutated (mutants 1, 2 and mutants 3, 4) and transfected into the A549 cells. Three base pairs of human MUC5AC regulatory regions (base pairs -971 to -939, mutant 1; -973 to -941, mutant 2; -235 to -203, mutant 3; -237 to -205; mutant 4) were mutated, as in table 1. Luciferase activity was measured from the control A549 cells and from each mutant-transfected A549 cell during the resting state and after stimulation with NE. Results shown are the average of four replicates, with standard deviation indicated by error bars (*, p<0.05 with respect to controls).

50 nM of neutrophil elastase markedly increased the transcriptional activity in transfectant A549 cells (Figure 1).

AG1478, PD98059, and CAPE all inhibited the transcriptional activity of MUC5AC promoter
To determine whether the EGF-R–MAP kinase-NF-κB pathway is involved in the NE–induced transcriptional activity of MUC5AC promoter, we added AG1478, a selective tyrosine kinase inhibitor; PD98059, a selective MEK inhibitor; and CAPE, an NF-κB inhibitor for 30 minutes prior to treatment with NE. These inhibitors completely suppressed the neutrophil elastase–induced transcriptional activity of the MUC5AC promoter (Figure 2).

Immunocytochemical staining
To determine whether NE increases the secretion of MUC5AC mucin from A549 cells, we performed immunocytochemical staining using the mouse monoclonal Ab to MUC5AC (clone 45M1, 1:1,000 New Markers, Fremont, CA). NE (50 nM) increased the secretion of MUC5AC mucin from A549 cells (data not shown).

Site-directed mutagenesis
Three base pairs of human MUC5AC regulatory regions which were known to be NF-κB binding sites were mutated (base pairs -971 to -939, mutant 1; -973 to -941, mutant 2: -235 to -203, mutant 3: -237 to -205, mutant 4) and transfected to the A549 cells. Except in mutant 3, NE (50 nM) did not increase the luciferase activity of the mutated promoter transfectants (Figure 3).

RT-PCR
To confirm the expression of MUC5AC mRNA, RNA analysis was carried out on the A549 cells using RT-PCR. NE (50 nM) increased the expression of MUC5AC mRNA (Figure 4). Signal transduction inhibitors, such as AG1478, PD98059, and CAPE, all suppressed the expression of MUC5AC mRNA by NE.

ERK1/2 were required for the activation of MUC5AC promoter by neutrophil elastase
To assess whether MAPK is involved in the NE–induced secretion of MUC5AC mucin in A549 cells, we performed immunoblotting with anti-phospho ERK1/2, anti-phospho P38, and anti-phospho JNK antibodies after stimulating the A549 cells with NE (50 and 100 nM). Only p-ERK1/2 developed; neither p38 nor JNK developed after stimulation with NE (Figure 5). The expressions of p-ERK1/2 after NE stimulation were completely suppressed after treatment with AG1478, PD98059, and CAPE.

NF-κB is involved in the induction of MUC5AC promoter activity by neutrophil elastase
Next, we assessed the effects of neutrophil elastase on the
DISCUSSION

In this study, we found that HNE increased the transcriptional activity of MUC5AC promoter and MUC5AC mRNA expression through the EGF–R ERK NF–κB pathway in A549 cells. Because the concentrations of HNE we used are within the range of airway concentrations of bronchial asthma patients, the increased mucus secretion of patients with asthma or COPD might result from the presence of NE in the airways. Neutrophils contain elastase at high concentrations (5 mM) in azurophil granules, and possess many biological properties, including pathogen killing, elastin fiber degradation, and MUC5AC mucin induction. Elastase has been reported to be a more potent secretagogue than other neutrophil proteases and secretory agonists (e.g., cathepsin G and histamine). However, until now, the mechanism which links mucin overproduction to HNE has not been well-defined. The A549 cell used in this study is the lung carcinoma cell line, which expressed high levels of MUC5AC mRNA and also possesses some of the characteristics of human bronchial epithelial cells. We hypothesized that HNE-induced mucin production is mediated through the EGF–R–MAP kinase pathway. To confirm this hypothesis, we transfected the A549 cells with the MUC5AC reporter plasmid (from C. Basbaum) and measured the luciferase activity after stimulation with HNE for 30 minutes. The most potent concentration for increasing the MUC5AC promoter activity was 50 nM of HNE. This increased transcriptional activity was markedly inhibited by pretreatment with EGF–R tyrosine kinase inhibitor (tyrphostin AG1478), MEK inhibitor (PD98059), or NF–κB inhibitor (CAPE). The MEK–MAPK transduction pathway was known to be downstream of EGF–R activation, and a selective inhibitor of MEK (PD98059) was reported to inhibit the activation of NF–κB in transfected A549 cells. EMSA showed increased nuclear NF–κB binding with the kB oligonucleotide probe by NE (Figure 6). This binding specificity was confirmed by competition and supershift assays.
production of MUC5AC in NCI-H292 cells. EGF-R is a member of the receptor tyrosine kinase superfamily and is involved in the regulation of proliferation and differentiation, primarily of epithelial cell types. The stimulation of EGF-R by its ligands, EGF, and transforming growth factor-α (TGF-α) causes MUC5AC mucin expression in airway epithelial cells. The importance of EGF-R in the production of MUC5AC mucin was demonstrated by the finding that HNE induced MUC5AC mucin production via the proteolytic activation of an EGF-R signaling cascade involving TGF-α. In the airways of rats, agarose plugs induce profound goblet cell metaplasia by causing the expression and activation of EGF-R. Activation of EGF-R is followed by the stimulation of various signaling pathways, including MAPKs p38, c-jun N-terminal kinase (JNK), ERK1/2, and big MAPK (BMK, ERK5), which subsequently results in the activation of various transcription factors (e.g., activator protein-1 and nuclear factor-κB), as well as gene transcription. Among the MAPKs, the three interconnected, relatively well-described pathways are the JNK, ERK, and p38 pathways. Each cascade is composed of at least three enzymes which are activated in series. The ERK1/2 (p44 and p42) pathways are the best characterized of the group. Selective p38 activation, however, was found to be important in the development of human non-small cell lung cancers. Because the MEK–MAPK transduction pathway is known to be downstream of EGF-R activation and because a selective inhibitor of MEK, PD98059, was reported to inhibit the MUC2 gene expression induced by Pseudomonas aeruginosa, we attempted to uncover the downstream pathway after EGF-R activation by NE. In our experiments, phosphorylated ERK1/2 were detected after NE treatment, but neither phosphorylated p38 nor phosphorylated JNK were detected by Western blotting. In inhibitor assays, AG1478, PD98059, and CAPE inhibited the development of phosphorylated ERK1/2. The mechanism of inhibition in the development of phosphorylated ERK1/2 by the NF-κB inhibitor, CAPE, is unknown, but we theorized that it might be due to the negative feedback mechanism. Here, we also showed that immunocytochemical staining of the A549 cells demonstrates the presence of MUC5AC mucin in the cytoplasm after HNE stimulation (data not shown). Inflammatory mediators from neutrophils, such as neutrophil elastase, may be present in chronically-diseased airways for prolonged periods of time, and it is believed that this chronic inflammation contributes to the profound airway remodeling observed in patients with long-term respiratory illnesses. For example, long-term exposure of normal human bronchial epithelial (NHBE) cells to IL-13 increases the percentage of Alcian blue/PAS-positive, mucus-producing cells in these cultures. This phenomenon was mediated by TGF-α, a potent ligand for EGF-R, which was rapidly released from NHBE cells in response to IL-13.

RT-PCR analysis also showed that 50 nM of HNE increased the expression of MUC5AC mRNA; it also showed that this increase was inhibited by pretreatment with CAPE, AG1478, or PD98059. Because CAPE is an inhibitor of NF-κB translocation, NF-κB is likely to be important to the expression of MUC5AC mRNA which is caused by HNE stimulation in A549 cells. NF-κB is a transcription factor which is critical for maximal expression of many cytokines that are involved in the pathogenesis of inflammatory diseases, such as adult respiratory distress syndrome (ARDS) and sepsis syndrome. NF-κB is activated by a variety of cytokines and mitogens. It is a key regulator of many of the genes which are involved in immune and inflammatory responses. Activation of NF-κB via Src-dependent dependent MAPK-p90rsk is important for Pseudomonas-induced mucin production in epithelial cells. The NF-κB family of transcription factors consists of homo- or hetero-dimeric subunits of the Rel family, including p65, p50, p52, c-Rel, and Rel-B. The regions between −973 to −939 and −237 to −203 of the MUC5AC promoter are known to be responsible for the transcriptional activation induced by NF-κB. The promoter regions including the specific NF-κB binding site in the human MUC5AC gene were cloned, and these transcriptional sites were upregulated by bacterial exoproducts. When these NF-κB binding sites were mutated and then transfected to the A549 cells in this experiment, the transcriptional activity was markedly inhibited in spite of HNE stimulation. These results show that the regions of the MUC5AC promoter between −973 to −939 and −237 to −203 are necessary for both basal promoter activity and NF-κB-induced transcriptional activation. Furthermore, EMSA using an oligonucleotide containing the previously identified MUC5AC NF-κB site demonstrated that nuclear extracts from HNE-treated A549 cells gave more intense bands than did nuclear extracts from untreated cells. For supershift assays, antibodies specific for p50, p65, or the c-Rel subunits of NF-κB were preincubated with nuclear extracts for 30 minutes on ice. The addition of p50, p65, and c-Rel antibody caused a decreased band intensity, which indicated a possible interaction. Thus, it appears that p50, p65, and c-Rel are all involved in the upregulation of MUC5AC by HNE. The specificity of protein binding to radio-labeled oligonucleotide was demonstrated by the addition of a 100-fold excess of unlabeled oligonucleotide. These results demonstrated that the NF-κB transcription factor activates MUC5AC expression through the NF-κB binding sites (−973 to −939 and −237 to −203) in the MUC5AC gene by binding as the p50/p65/c-Rel heterodimer.

It has been found that mucus hypersecretion can be triggered by a variety of factors, including proinflammatory cytokines (e.g., IL-4, IL-9, IL-13), EGF, TNF-α, ATP, bacterial exoproducts, and neutrophils. Mucus is essential because of its...
role in protecting the airways. In chronic airway disease, however, mucus hypersecretion is an important factor in morbidity and mortality. NE and oxidant radical are especially important in the pathogenesis of pulmonary emphysema. NE increases the reactive oxygen radicals in lung cells, which contribute to cell death. NE is also known to increase the MUC5AC mRNA levels by enhancing mRNA stability or by inducing reactive oxygen species. MAPKs may be directly activated by oxidants such as H$_2$O$_2$. Oxidative stress causes the synthesis of MUC5AC mucin in NCI-H292 cells by transactivation of p44/42MAPK and EGF–R. EGF–R serves a central role as a primary regulator of epithelial function, transducing extracellular signals from its activating ligands into intracellular signaling cascades through dimerization and transphosphorylation catalyzed by the intrinsic tyrosine kinase. Depending on the triggering stimuli, the signal transduction pathways which control mucin transcription have distinct pathways. NE also promotes the detachment of the epithelial cells of the airway and changes the morphology of bronchial epithelial cells through the activation of MAP kinase.

In this experiment, we defined the role of NE in MUC5AC mucin secretion and determined its signal transduction pathways in A549 cells. The MUC5AC mucin synthesis might be caused by both the direct stimulation of NE and the oxidant radicals which were produced from the A549 cells by NE. We speculate that if the signal is mediated by oxidative stress, such as the activation of ERK1/2 by H$_2$O$_2$, antioxidants such as N-acetylcysteine will be effective in reducing MUC5AC mucin synthesis. We hope that this study will provide potential therapeutic targets in neutrophil-predominant airway inflammation and mucus obstruction of the airways.

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