

Staphylococcus aureus Enterotoxin B Regulates Prostaglandin E₂ Synthesis, Growth, and Migration in Nasal Tissue Fibroblasts

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Background. Superantigens and eicosanoids are important amplifiers and regulators of inflammation in airway diseases. We therefore studied the possible influence of *Staphylococcus aureus* enterotoxin B (SEB) on the cyclooxygenase (COX) pathway and basic functions of airway structural cells.

Methods. Fibroblasts were isolated from nasal inferior turbinate tissue and cultured in the presence of different concentrations of SEB. Preincubation with interferon (IFN)- γ was performed to induce expression of major histocompatibility complex (MHC) class II receptors. Prostaglandin E₂ (PGE₂) production was assayed by enzyme-linked immunosorbent assay, and levels of COX-2 and prostanoid E receptors 1–4 (EP_{1–4}) were assayed by real-time polymerase chain reaction. Migration and growth tests were performed, and SEB was localized within the cells by confocal microscopy.

Results. Stimulation with IFN- γ and SEB significantly down-regulated PGE₂, COX-2, and EP₂ expression but not EP₁, EP₃, or EP₄ expression. The enterotoxin blocked cell growth but increased the fibroblast migration rate. SEB was localized within the cell in the presence and absence of MHC-II, suggesting that mechanisms other than conventional binding may allow the enterotoxin to enter the cell.

Conclusions. These findings may have major implications for our understanding of the role played by bacterial superantigens in regulating the inflammatory and remodeling mechanisms of upper airway diseases and hence may help elucidate the pathophysiology of these diseases.

Fibroblasts are important sentinel cells of the immune system, playing a critical role in chronic inflammation and tissue repair. However, the implication of these cells in the regulation of immune responses has long been neglected. Recently, several studies have demonstrated that human lung fibroblasts activated through the CD40-CD40L bridge may up-regulate prostaglandin E₂ (PGE₂) production [1] and, hence, hematopoietic cells infiltration in damaged tissue [2]. This interaction activates the transcription factor NF- κ B, resulting in the

synthesis of high levels of cytokine mediators (e.g., interleukin [IL]-6 and IL-8), adhesion molecules (e.g., intercellular and vascular cell adhesion molecules), and cyclooxygenase (COX) metabolites [3]. In the airways, fibroblasts play a crucial role in the remodeling process. These cells are a primary cellular source of collagens and growth factors, which play a crucial role in epithelial regeneration and tissue remodeling processes [4].

The potential role played by superantigens in the pathogenesis of airway diseases has become a relevant issue during the last years [5]. Furthermore, it is known that prostanoids, metabolites of arachidonic acid, are crucial molecules in the control of inflammatory and remodeling processes in airways [6, 7]. We have previously demonstrated that immune response to *Staphylococcus aureus*, in terms of IgE specific to *S. aureus* enterotoxins, correlates with total IgE levels and severity of inflammation in patients with nasal polyposis [8, 9]. Additionally, this immune response to *S. aureus* enterotoxins was associated with an amplification of the eosinophilic inflammation

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and an up-regulation of leukotrienes and lipoxins within nasal polyp tissue [9], which is also characterized by reduced PGE₂ concentrations [7].

Additionally, *in vitro* experiments have shown that *S. aureus* enterotoxins can regulate the secretion of interstitial collagenase and stromelysin in human fibroblastlike synoviocytes [10, 11], and these cells can function as accessory cells for enterotoxin-mediated T cell activation [12]. More recently, it was demonstrated that *S. aureus* enterotoxins can induce IL-6 and IL-8 gene expression on human fibroblastlike synoviocytes isolated from patients with rheumatoid arthritis [13]. Production of PGE₂ can also be enhanced by these superantigens in bovine mononuclear cells [14]. Until now there have been no studies regarding the role played by *S. aureus* superantigens on structural cells from the upper airways. Given the previous findings, we aimed to study the capacity of *S. aureus* enterotoxin B (SEB) in regulating the COX pathway and vital cell processes, such as growth and migration, in nasal tissue fibroblasts.

MATERIALS AND METHODS

Samples. Fibroblasts were isolated from inferior turbinate tissue obtained from patients ($n = 5$) who were undergoing septoplasty or rhinoseptoplasty because of anatomical variations and were not suffering from any sinus disease. Surgery was performed in the Department of Otorhinolaryngology at Ghent University Hospital. All patients gave informed consent before their participation, and the study was approved by the ethical committee of Ghent University Hospital.

Reagents. Dulbecco's PBS, penicillin-streptomycin (penicillin, 5000 IU/mL; streptomycin, 5000 µg/mL), and trypan blue (0.4% solution in PBS) were obtained from Invitrogen. Minimum essential medium (MEM), Opti-MEM I reduced serum medium, L-glutamine (200 mmol/L), trypsin-EDTA (1×), and fetal bovine serum (FBS) (qualified; origin, United States) were purchased from Gibco (Life Technologies). Ultrosor G (5%) was obtained from Biospera. Recombinant human interferon (IFN)-γ was obtained from R&D Systems, and SEB was obtained from Sigma Chemicals. SYBR Green I Master mix, Aurum Total RNA, and Script cDNA synthesis kits were obtained from Bio-Rad Laboratories.

Fibroblast isolation from nasal tissue. Nasal tissues obtained during surgical operations were rinsed several times with Opti-MEM I supplemented with 5% FBS, 5% Ultrosor G, 2 mmol/L glutamine, and penicillin (50 IU/mL)-streptomycin (50 µg/mL) and cut into small pieces (~1 mm²). Diced specimens were then plated (density, 9 pieces/6-well tissue culture dish) and incubated in a humidified atmosphere containing 5% CO₂ at 37°C, until a monolayer of fibroblastlike cells was observed to be confluent. Then the explanted tissues were removed, and cells were trypsinized and replated into 250-cm² tissue culture Falcon tubes at a final volume of 5 mL. The me-

dium was changed every 3 days for 2–3 weeks until 90% confluence was obtained. Subsequently, the cells were split and passaged.

Cell stimulation. Cells were stimulated with 0.01, 0.1, or 1 µg/mL SEB or 10 ng/mL IL-1β for 24 h at 37°C in 5% CO₂. To induce receptors for major histocompatibility complex (MHC) class II, cells were pretreated with IFN-γ (10 ng/mL) for 24 h at 37°C in 5% CO₂ and were then stimulated with 0.01, 0.1, or 1 µg/mL SEB for 24 h at 37°C and 5% CO₂. For each sample in each experiment, we used a negative control that consisted of cells incubated only with culture medium (MEM).

The cells were characterized with antibodies against vimentin, cytokeratin, and α-smooth muscle actin, using the peroxidase-anti-peroxidase technique. Stimulated cell cultures were centrifuged at 4°C, and the supernatants were collected and stored at -20°C until use. Subsequently, cells were trypsinized, washed, resuspended in Opti-MEM I, and then centrifuged. After the supernatants were removed, a portion of the cells were resuspended in lysis buffer for posterior RNA extraction, and another portion was stored at -80°C.

Enzymatic immunoassays. Concentration of PGE₂ were measured by ELISAs in cell culture supernatants according to the manufacturer's instructions (Oxford Biomedical); the intra- and interassay coefficient of variation for all ELISAs was <10%.

Quantitative real-time polymerase chain reaction (PCR). Cell cultures were centrifuged, and pellets containing 1×10^6 cells were disrupted and homogenized in lysis solution. Total RNA was extracted using the Aurum Total RNA Mini Kit, and cDNA was synthesized from 1 µg of RNA using the Script cDNA Kit, in accordance with the manufacturer's instructions. Amplification reactions were performed with the iCycler iQ Real-Time PCR Detection System, with the specific primer sets (table 1). PCRs included 20 ng of cDNA from cultured cells, 1× SYBR Green I Master mix, and 300 nmol/L each primer. The PCR protocol consisted of 1 cycle at 95°C for 10 min followed by 40 cycles at 95°C for 30 s and at 62°C for 1 min. Ubiquitin C and hypoxanthine phosphoribosyltransferase 1 were used as internal control genes for normalization. Primer sequences for these genes were obtained from the Real-Time PCR Primer and Probe Database of the Department of Medical Genetics, University of Ghent (<http://medgen.ugent.be/rtprimerdb>). The relative number of molecules for each gene was expressed in relative expression units quantified per 20 ng of cDNA sample and was determined by the ΔΔCt value method.

Cell growth assay. Confluent cells in 25-cm² Falcon dishes were trypsinized with preheated trypsin-EDTA and resuspended in 5 mL of Opti-MEM I. Living cells were counted with a Neubauer counting chamber, seeded (10,000 cells/well) in a 24-well plate, and incubated with Opti-MEM I, IFN-γ (10 ng/mL), or SEB (0.1 µg/mL) at 37°C and 5% CO₂ for 24, 48, and 72 h. At each time point, cells were again trypsinized with preheated trypsin-EDTA and resuspended in 1 mL of Opti-MEM I, and the

Table 1. Primer sequences used for real-time polymerase chain reaction amplification.

Gene	Primer (5'→3')		Amplicon size, bp	GenBank accession no.
	Forward	Reverse		
HLA-DR1 α	CATCAAGGGAGTGCGCA	CTCCATGTGCCTTACAGAGGC	64	NM_019111
COX-2	GCTGGAACATGGAATTACCCA	CTTTCTGTACTGCGGGTGGAA	88	XM_051899
EP ₁	GATGGTGGGCCAGCTTGTC	GCCACCAACACCAGCATTG	73	NM_000955
EP ₂	GACCGCTTACCTGCAGCTGTAC	TGAAGTTGCAGGCGAGCA	73	NM_000956
EP ₃	AAGGCCACGGCATCTCAGT	TGATCCCCATAAGCTGAATGG	76	NM_000957
EP ₄	ACGCCGCCTACTCTACATG	AGAGGACGGTGGCGAGAAT	63	NM_000958

NOTE. COX-2, cyclooxygenase 2; EP₁₋₄, prostanoid E receptors 1–4.

number of living cells at each time point was quantified with trypan blue using the Neubauer counting chamber.

Cell migration assay. Fibroblasts were seeded in a 35-mm-diameter 6-well plate, incubated at 1×10^5 cells/well in serum-free MEM, and allowed to grow until confluency for 7 days. Then the medium was replaced by $1 \times$ Dulbecco's PBS, and cells were incubated at 37°C in 5% CO₂ for 5 min. The monolayer was wounded by pressing a sterile razor blade down onto the well, making a sharp visible demarcation at the wound edge. The blade was then gently moved to one side to remove a part of the cell monolayer. Two separated 15–20-mm wounds were made in the same well. The wounded layers were washed 2 times with Opti-MEM to remove the cell debris and incubated with Opti-MEM I, IL-1 β (10 ng/mL), IFN- γ (10 ng/mL), or SEB (0.1 μ g/mL) at 37°C and 5% CO₂. Migrating cells reaching a line (L2) situated 240 μ m from the culture wound after 24 (L2D1) and 48 h (L2D2) were counted using a counting eyepiece with grid graticule (10 \times 10 mm). The first line of the grids falls together with the sharp beginning demarcation of the wound. Ten places were counted in each wound.

Cellular localization of SEB. Cellular localization of SEB was assayed by immunofluorescence confocal microscopy. Nasal tissue fibroblasts were cultured with 0.1 μ g/mL SEB, with or without IFN- γ (10 ng/mL), on poly-D-lysine-coated glass slides for 24 h at 37°C in 5% CO₂. The slides were then air dried for 24 h at room temperature and stored at -20°C. At the time of staining they were fixed in acetone for 10 min at 4°C and blocked using 0.4% fish-skin gelatine/1 \times PBS for 2 h at room temperature. The cells were then incubated with rabbit polyclonal antibody anti-SEB (1:500; Sigma) for 1 h and with the secondary antibody anti-rabbit Alexa Fluor 526 (1:2000; Molecular Probes) for 1 h at room temperature; both primary and secondary antibodies were diluted in 0.4% fish-skin gelatine/1 \times PBS. After nuclear staining, 4',6-diamidino-2-phenylindole (DAPI; Molecular Probes) was added for 5 min, and the slides were mounted in Vectashield mounting medium (Vector Laboratories). Cellular localization of the enterotoxin was evaluated using a Bio-Rad Radiance 2100 laser scanning confocal microscope system with a 4-line argon primary laser, a green helium-neon laser, and a blue laser diode. Image analysis was performed using Adobe Photo-

shop (version 7.0) and Confocal Assistant (version 4.02, Todd Clark Brelje) software programs.

Statistical analysis. All data were analyzed using MedCalc software (version 6.0) and are presented as medians and interquartile ranges. Data were compared within different patient subgroups by use of the Kruskal-Wallis test (*H* test). The Wilcoxon test for unpaired samples (or Mann-Whitney *U* test) was applied to evaluate the statistical differences between patient groups. Our criterion for significance was $P \leq .05$.

RESULTS

Effect of SEB on PGE₂ production. Nasal tissue fibroblasts cultured in MEM in the absence of stimulus produced significant baseline concentrations of mRNA of COX-2 and PGE₂. Stimulation with SEB without preincubation with IFN- γ did not induce any change in PGE₂ production or COX-2 mRNA expression. Induction of MHC-II molecule expression in fibroblasts by stimulation with IFN- γ was verified by real-time PCR, demonstrating HLA-DR1 α gene expression in all IFN- γ -stimulated cells compared with nonstimulated cells (figure 1). Furthermore, preincubation with IFN- γ led to a significant decrease

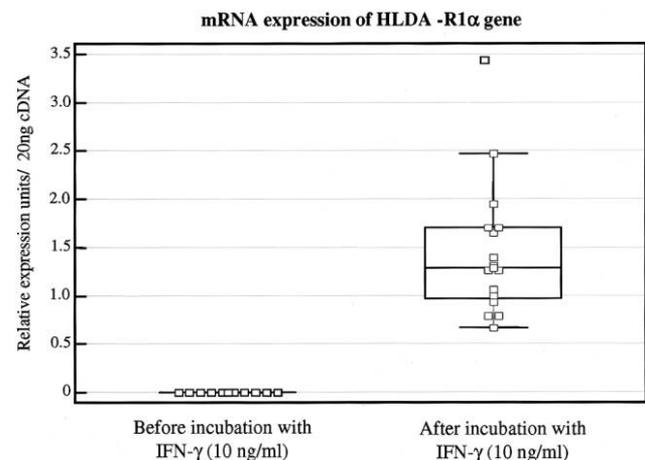


Figure 1. Box-and-whisker plots representing the expression levels of the HLA-DR1 α gene in nasal tissue fibroblasts before and after incubation with interferon (IFN)- γ (10 ng/mL).

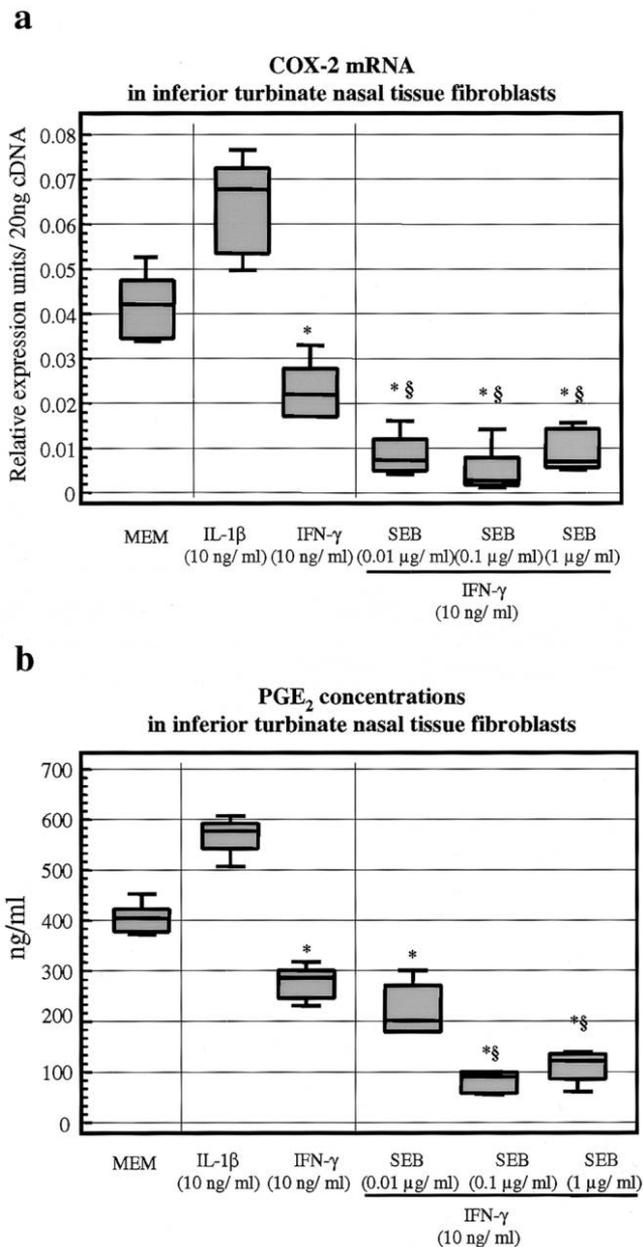


Figure 2. mRNA levels of cyclooxygenase (COX)-2 (*a*) and concentrations of prostaglandin E₂ (PGE₂) (*b*) in nasal tissue fibroblasts after stimulation with interleukin (IL)-1β (10 ng/mL), interferon (IFN)-γ (10 ng/mL), or *Staphylococcus aureus* enterotoxin B (SEB; 0.01, 0.1 or 1 μg/mL). *Significant difference compared with cells incubated with minimum essential medium (MEM); §significant difference compared with cells stimulated with IFN-γ (10 ng/mL) ($P < .05$, Mann-Whitney *U* test for independent samples).

in COX-2 mRNA (figure 2*a*) and endogenous PGE₂ (figure 2*b*) compared with nonstimulated cells. Interestingly, this down-regulation was amplified after incubation with SEB at 0.1 and 1 μg/mL in a dose-independent manner. In addition, incubation with IL-1β significantly increased the expression of COX-2 and PGE₂ release. Furthermore, expression of the transcripts expressing the genes encoding for prostanoid E receptors (EP₁,

EP₂, EP₃, and EP₄) was analyzed. Real-time PCR results showed no expression of EP₄ receptor in nasal fibroblasts (data not shown). SEB did not induce any change in the expression of EP₁ (figure 3*a*) and EP₃ (figure 3*c*) receptors; however, the enterotoxin down-regulated the EP₂ receptor after preincubation with IFN-γ, and this effect was dose dependent (figure 3*b*).

Effect of SEB on growth and migration. Results of the growth test were expressed as numbers of living cells after 24, 48, and 72 h. The number of cells per stimulus was compared with the number of cells cultured only in MEM, which were considered to be controls for normal growth. Cells in MEM showed a significant increase in proliferation after 72 h. The same pattern was observed in cells stimulated with IFN-γ or IL-1β (figure 4). When cells were incubated with SEB, however, cell proliferation was inhibited. This process was IFN-γ independent and was counterbalanced by coexposure with the cytokine. Furthermore, the cell migration capacity was not changed by incubation with IFN-γ or SEB. After preincubation with IFN-γ, however, SEB significantly induced this process (figure 5).

Cellular localization of SEB. Cellular localization of SEB was assayed by confocal microscopy. Confocal images of nasal fibroblasts after incubation with 0.1 μg/mL SEB for 24 h showed that the enterotoxin was located mainly in the cytoplasm but also partly in the nucleus of the cell (figure 6*c* and 6*d*), indicating spontaneous penetration into the cell. A different distribution was observed after preincubation with IFN-γ followed by stimulation with 0.1 μg/mL SEB (figure 6*e* and 6*f*). The enterotoxin was located mainly in the periphery of the cell cytoplasm and around the nucleus, suggesting the binding of SEB to MHC-II molecules on the cell surface.

DISCUSSION

Eicosanoid production is differentially regulated and correlates with the degree of inflammation in patients with nasal polyps with and without an immune response (specific IgE antibodies) to *S. aureus* enterotoxins [9]. PGE₂ plays an important role in the regulation of tissue repair events during airway diseases, and this mechanism may be influenced by triggers such as viral or bacterial infections. Therefore, we aimed to investigate whether staphylococcal enterotoxins may affect the upper airway remodeling process and influence the eicosanoid pathway of nasal tissue fibroblasts.

In this study, we were able to demonstrate that the levels of endogenous PGE₂, COX-2 mRNA, and EP₂ receptor significantly decreased after stimulation with SEB in IFN-γ-pretreated cells but not in cells cultured without the cytokine, suggesting that this effect may be mediated by cross-linking with the MHC-II receptor. Incubation with IFN-γ was performed to induce the MHC-II molecules, which are normally not expressed in fibroblasts, and mimic the classic binding of superantigens to the cell. This cytokine has been demonstrated to induce the syn-

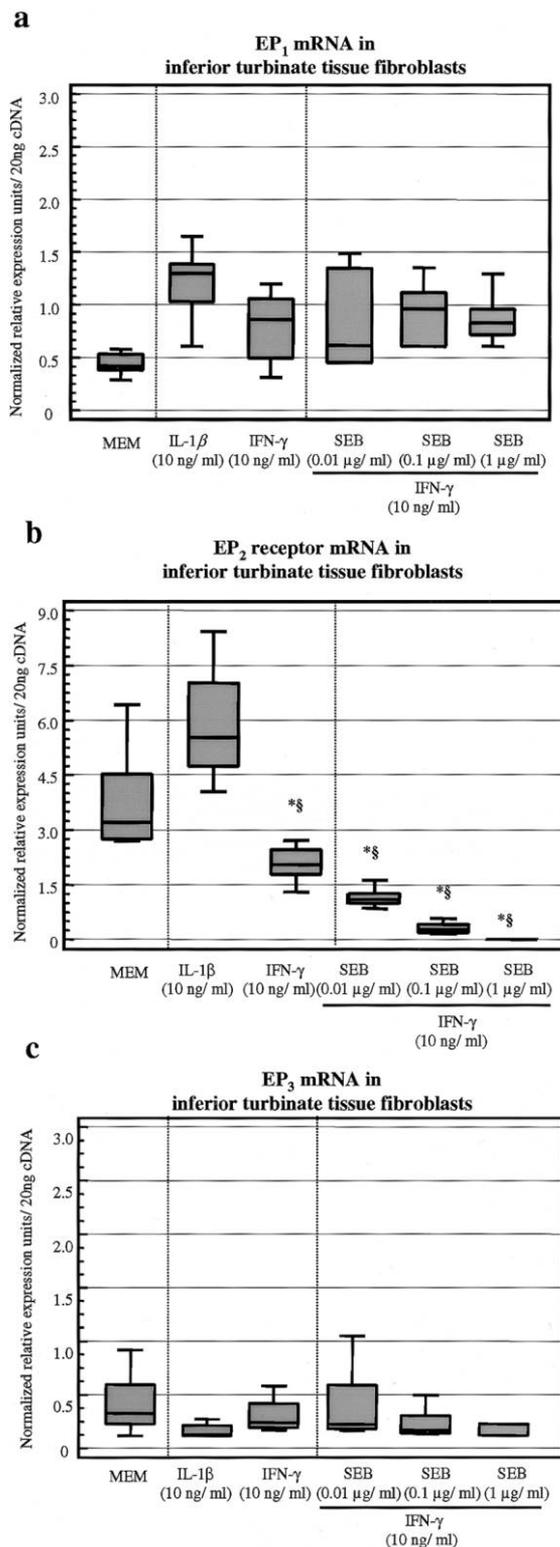


Figure 3. mRNA levels of prostanoid E (EP) receptors EP₁ (a), EP₂ (b), and EP₃ (c) in nasal tissue fibroblasts after stimulation with interleukin (IL)-1 β (10 ng/mL), interferon (IFN)- γ (10 ng/mL), or *Staphylococcus aureus* enterotoxin B (SEB; 0.01, 0.1, or 1 μ g/mL). *Significant difference compared with cells incubated with minimum essential medium (MEM); §Significant difference compared with cells stimulated with IFN- γ (10 ng/mL) ($P < .05$, Mann-Whitney U test for independent samples).

thesis and activation of cytosolic phospholipase A₂ (cPLA₂) protein in a human bronchial epithelial cell line [15]. However, it seems that nasal fibroblasts may react differently to the cytokine, because COX-2 and PGE₂ were significantly down-regulated compared with nonstimulated cells. Additionally, staphylococcal enterotoxins have been demonstrated to induce PGE₂ production and COX-2 and cPLA₂ expression in IFN- γ -treated fibroblastlike synoviocytes [10]. However, in nasal fibroblasts, SEB acted as an agonist of the cytokine by accentuating the blockage of the COX pathway.

There have been no previous studies showing the effect of superantigens on PGE₂ release in nasal fibroblasts. However, Clement et al. have demonstrated highly focalized intracellular reservoirs of *S. aureus* (including the nasal epithelium, glandular tissue, and myofibroblasts) in endonasal mucosa from patients with recurrent *S. aureus* rhinosinusitis [16]. These findings support our thinking that the presence of SEB in nasal tissue may exacerbate airway inflammation by down-regulating the production of anti-inflammatory mediators, such as PGE₂.

Moreover, staphylococcal superantigens may act as potent leukocyte and neutrophil chemoattractants [17]. In support of these findings, we have found that SEB significantly induced fibroblast migration. This effect seems to be mediated by the binding of MHC-II, because no effect was observed in the presence of the enterotoxin or IFN- γ alone. IFN- γ has been reported to increase the expression of both intercellular adhesion molecule 1 and vascular cell adhesion molecule in human lung fibroblasts [18], important regulators of cell adhesion and migration processes. Furthermore, it has been shown that SEB can induce T cell adhesion to endothelial cells (pretreated with IFN- γ) in an MHC-II-dependent mechanism [19], which may confirm the findings of this study. However, although it was not our current objective, it may be useful to evaluate the expression of cell migration-related molecules to learn more about the mechanisms modulating this process.

SEB blocked proliferation in the cell culture when it was applied in the absence of IFN- γ , and this effect may be restored by the cytokine. Although the effect of SEB on cell proliferation is unknown, the superantigen may block this process by cross-linking receptors other than MHC-II or simply by altering cell membrane permeability or activating molecules that regulate this cell function. Furthermore, PGE₂ has been reported as an important inhibitor of fibroblast proliferation and migration [20, 21]. However, analyzing our results, we suggest that the effect of SEB on fibroblast proliferation is not linked to PGE₂ regulation. On the contrary, the migration process seems to be influenced by the decrease in eicosanoid levels after SEB exposure. However, this can be confirmed only by performing experiments in the presence of PGE₂ and MHC-II antagonists, which should be the aim of further studies.

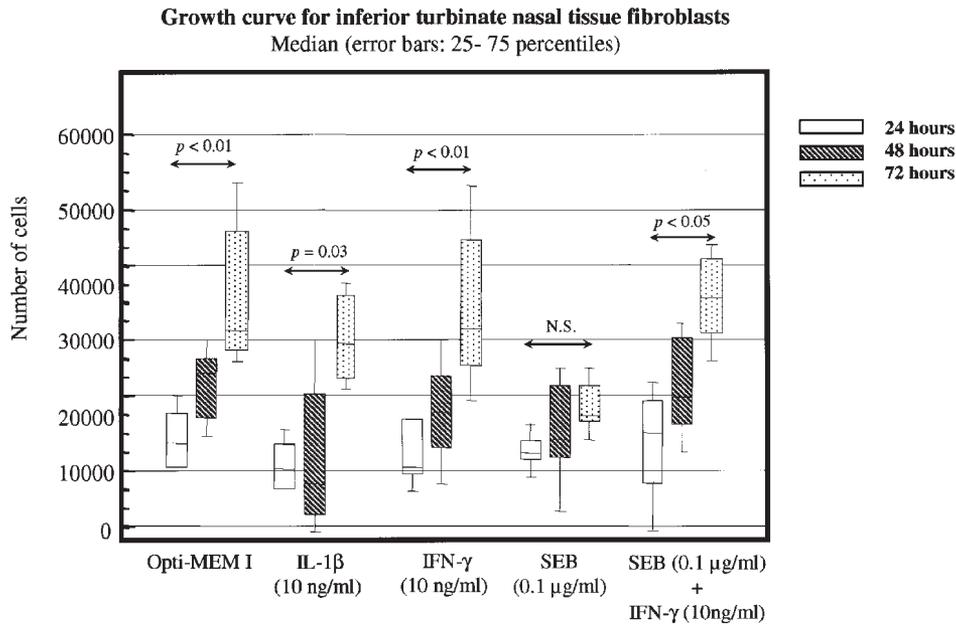


Figure 4. Growth curve for nasal tissue fibroblasts showing the number of proliferated cells after incubation with interleukin (IL)-1 β (10 ng/mL), interferon (IFN)- γ (10 ng/mL), or *Staphylococcus aureus* enterotoxin B (SEB; 0.1 μ g/mL) for 24, 48, and 72 h. *P* values were determined by the Mann-Whitney *U* test for independent samples. N.S., not significant.

Finally, confocal microscopy showed the presence of SEB within cells with and without MHC-II receptors. These findings suggest that the enterotoxin may enter the cell via alternate mechanisms or receptors in cells that lack MHC-II or T cell receptors. Studies have shown that SEB may cross the epithelium by facilitated transcytosis [22], and the

existence of alternative binding site proteins has been also reported in monkey kidney fibroblasts [23]. Our results indicate that the effect of SEB on human nasal fibroblast metabolism may depend on how the molecules enter the cell and may or may not depend on the classic superantigen-binding mechanism.

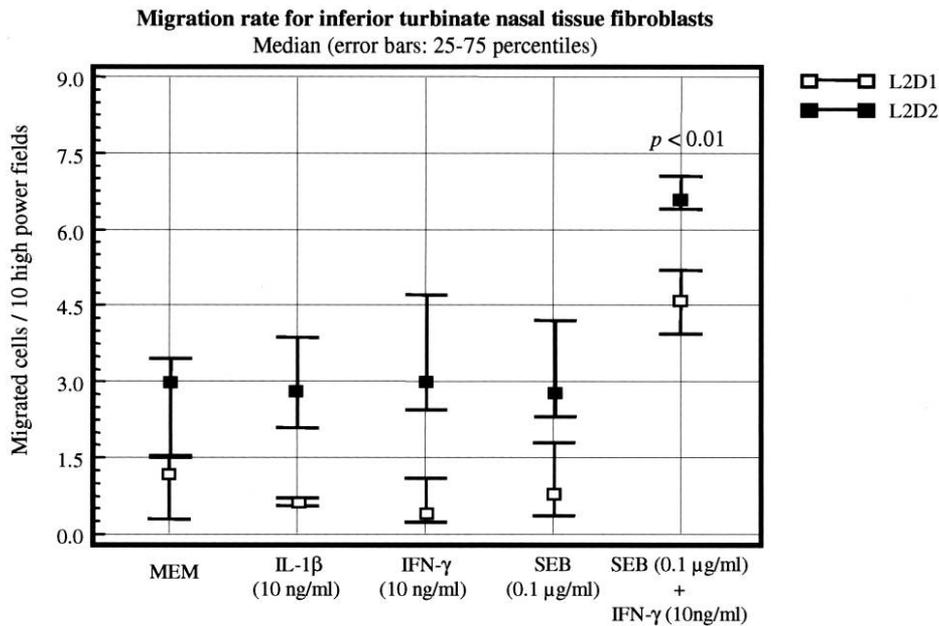


Figure 5. Migration test for nasal tissue fibroblasts after stimulation with interleukin (IL)-1 β (10 ng/mL), interferon (IFN)- γ (10 ng/mL), or *Staphylococcus aureus* enterotoxin B (SEB; 0.1 μ g/mL). Graph shows the number of cells migrating 240 μ m from the wound site after 24 h (L2D1) and 48 h (L2D2). MEM, minimum essential medium.

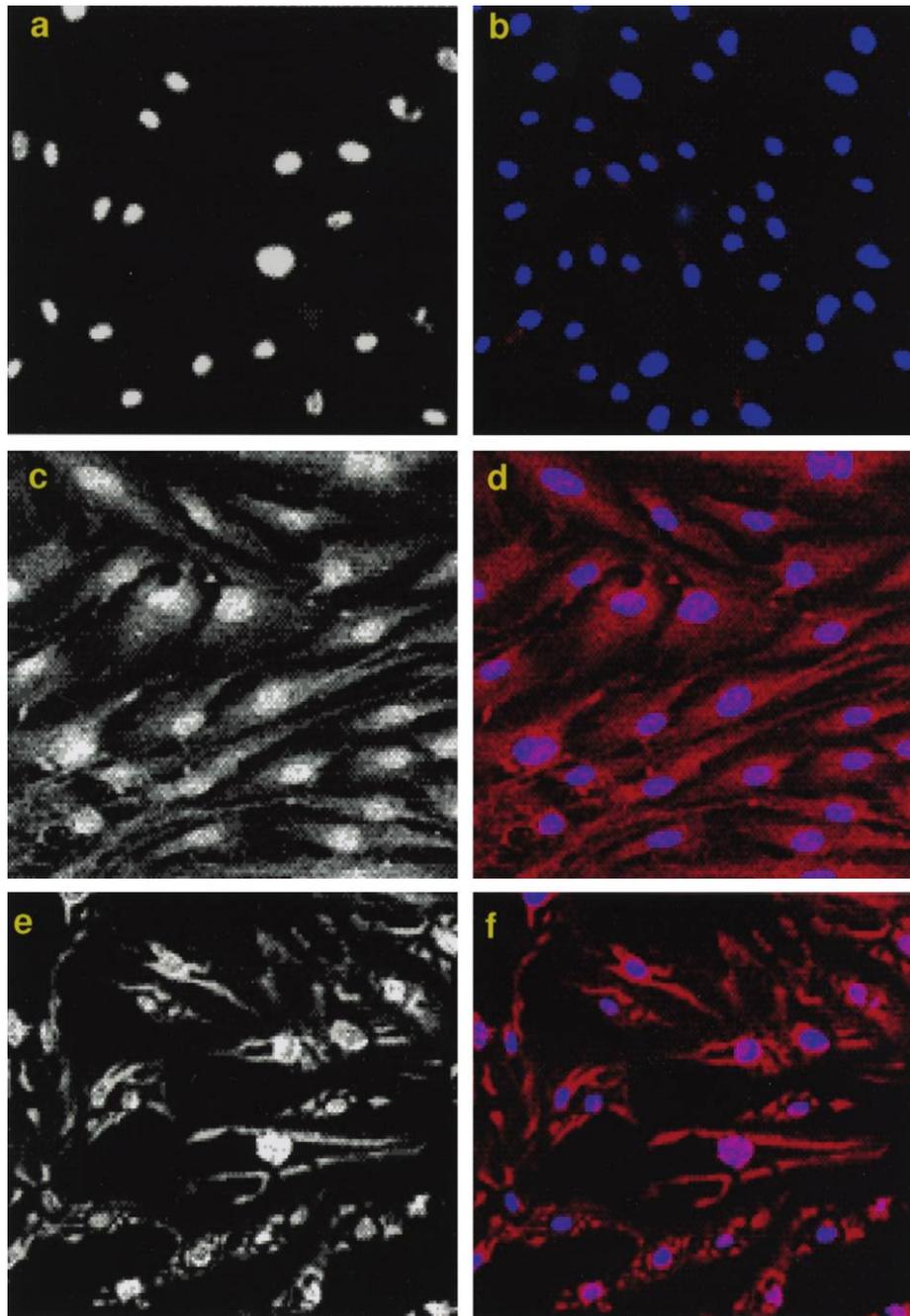


Figure 6. Confocal microscopic images (original magnification, $\times 40$) of nasal tissue fibroblasts incubated with minimum essential medium (*a* and *b*), stimulated with *Staphylococcus aureus* enterotoxin B (SEB; $0.1 \mu\text{g}/\text{mL}$) for 24 h (*c* and *d*), or preincubated with interferon- γ ($10 \text{ ng}/\text{mL}$) and stimulated with SEB ($0.1 \mu\text{g}/\text{mL}$) for 24 h (*e* and *f*). Images show that SEB is internalized in the nasal fibroblasts and can be localized in the cytoplasm and in the nucleus of the cell.

In conclusion, SEB may down-regulate COX-2 gene expression and PGE₂ production in human nasal tissue fibroblasts, and this action seems to involve EP₂ but not EP₁, EP₃, or EP₄ receptors. Additionally, this enterotoxin may affect important basic cell functions, such as growth and migration. Determining the mechanism behind this action was not the objective of this study, but it seems that mechanisms other than the conventional binding via MHC-II allow the enterotoxin to enter the

cell and influence its metabolism. The present study will be followed by experiments using PGE₂ and MHC-II inhibitors, which can provide more information on the regulatory mechanisms acting on these cells. Triggering of the arachidonic acid metabolism is one of the cellular events regulating the pro- and anti-inflammatory immune response in diseases. The findings of this study may have major implications for our understanding of the role played by bacterial superanti-

gens in regulating the inflammatory and remodeling mechanisms of upper airway diseases and, hence, help to elucidate their pathophysiology.

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