

Inhibition by glucocorticoids of the interleukin-1 β -enhanced expression of the mast cell growth factor SCF

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- 1 Stem cell factor (SCF) is a major mast cell growth factor that promotes differentiation and chemotaxis of mast cells and inhibits their apoptosis.
- 2 We evaluated the effect of interleukin (IL)-1 β , a major pro-inflammatory cytokine, on the constitutive expression of SCF and studied the effects of two glucocorticoids, budesonide and dexamethasone, on the IL-1 β -enhanced SCF expression.
- 3 Human lung fibroblasts in culture were serum-starved for 48 h and treated with IL-1 β , budesonide and/or RU486. SCF cDNA was quantified after total RNA reverse transcription by on-line fluorescent polymerase chain reaction. SCF protein was quantified by ELISA.
- 4 IL-1 β induced an increase in SCF mRNA (+91% at 2.5 h) and protein production (+32%) by human lung fibroblasts in culture ($P < 0.001$).
- 5 Budesonide inhibited IL-1 β -induced SCF mRNA expression (–68%) at 2.5 h and even more so at 10 h (–192%) ($P < 0.001$). The expression of SCF protein also decreased by 3.5-fold at 10 h. Results were similar with dexamethasone. The glucocorticoid antagonist RU486 cancelled the effects induced by the glucocorticoids.
- 6 Increased SCF mRNA levels were associated with increased stability of this mRNA as measured after treatment with actinomycin D (1.9-fold at 2.5 h). Budesonide decreased this IL-1 β -enhanced stability by about 1.5-fold ($P < 0.001$).
- 7 We conclude that in 'inflammatory' conditions, mimicked *in vitro* by IL-1 β , glucocorticoid treatment inhibits expression of the mast cell growth factor SCF. The reduced number and activation of mast cells observed in the bronchi of asthmatic patients treated by glucocorticoids may be due in part to this effect.

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Keywords: SCF; IL-1 β ; budesonide; dexamethasone; fibroblasts; mast cells; lung; asthma

Abbreviations: COX-2, cyclooxygenase type-2; DMEM, Dulbecco's modified eagle's medium; ELISA, enzyme-linked immunosorbent assay; FCS, foetal calf serum; GR, glucocorticoid-receptor; IL-1 β interleukin (IL)-1 β ; NF- κ B, nuclear factor-kappa B; PCR, polymerase chain reaction; RT, reverse transcription; SCF, stem cell factor

Introduction

Stem cell factor (SCF), also termed Kit ligand, steel factor, or mast cell growth factor (Huang *et al.*, 1990; Martin *et al.*, 1990; Zsebo *et al.*, 1990), acts as an important growth factor for human mast cells (Galli *et al.*, 1995). *In vitro* it induces the proliferation and differentiation of immature mast cells from bone marrow CD34+ progenitor cells (Rottem *et al.*, 1994). SCF also aids mast cell survival by inhibiting their apoptosis (Iemura *et al.*, 1994). Moreover, it is a potent chemotactic factor for mast cells (Nilsson *et al.*, 1994) and promotes their adhesion to the extracellular matrix (Dastyh & Metcalfe, 1994). In addition, *in vivo*, SCF induces mast cell hyperplasia after subcutaneous injection in humans (Costa *et al.*, 1996). Finally, *in vitro* it greatly enhances antigen-induced degranulation of human lung-derived mast cells

(Bischoff & Dahinden, 1992) as well as inducing mast cell degranulation itself alone *in vitro* (Takaishi *et al.*, 1994) and *in vivo* (Costa *et al.*, 1996). Therefore, SCF may be involved in many diseases associated with a local increase in the number and activation of mast cells, such as occurs in asthma (Ammit *et al.*, 1997).

In asthmatic patients, large quantities of the pro-inflammatory cytokine interleukin (IL)-1 β have been reported in bronchoalveolar lavage (Borish *et al.*, 1992). IL-1 β may be released by structural cells, such as epithelial cells (Devalia *et al.*, 1993) and fibroblasts (Pan *et al.*, 1996), as well as by infiltrated inflammatory cells, including eosinophils (Barnes, 1990) and mast cells (Moller *et al.*, 1998). IL-1 β is known to upregulate the expression of many genes involved in the inflammatory process, including RANTES (Saji *et al.*, 2000), eotaxin (Ghaffar *et al.*, 1999), IL-6 and IL-8 (Agarwal *et al.*, 1995) and the IL-1 β gene itself (Hiscott *et al.*, 1993). However, the effect of IL-1 β on SCF expression by pulmonary cells is not known.

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Glucocorticoids remain the major treatment for asthma. This treatment is associated with a reduced number of infiltrated mast cells in the bronchial mucosa of asthmatic patients (Jeffery *et al.*, 1992). Glucocorticoids affect cells in many ways, but their major therapeutic benefit is likely to occur through the regulation of the expression of genes involved in inflammation (for review, see Barnes & Adcock, 1993). In particular, as recently reported by Kassel *et al.* (1998), the regulation of the constitutive expression of SCF by glucocorticoids shows a time-dependent biphasic effect in human lung fibroblasts in culture.

Here we report our investigation of the regulation of SCF expression in inflammatory conditions mimicked by the addition of IL-1 β , and the effect of the glucocorticoids budesonide and dexamethasone on the IL-1 β -induced SCF expression by human pulmonary fibroblasts in culture.

Methods

Culture of human lung fibroblasts

Human lung derived fibroblasts were obtained by the explant technique, as previously described (Kassel *et al.*, 1998). Briefly, macroscopically normal human lung tissue was separated into fragments within 1 h of resection for bronchocarcinoma. Fibroblasts were cultured in Dulbecco's modified Eagle's medium (DMEM)/F-12 (1:1) (GIBCO BRL, Cergy Pontoise, France), supplemented with 10% foetal calf serum (FCS), penicillin and streptomycin, and incubated in a humidified mixture of 95% air and 5% CO₂ at 37°C. They were then replated in 25 cm² tissue culture flasks (Costar, Acton, MA, U.S.A.) and were characterized as fibroblasts morphologically and by immunocytochemistry with an antifibroblast monoclonal antibody (5B5) (DAKO S.A., Trappes, France). They subsequently were split 1:4 at confluence and passaged. Fibroblasts were used at passage 7.

Cell treatment

At confluence, fibroblasts were placed into a quiescent state by reducing the FCS content to 0.3% (starving medium) for 48 h. IL-1 β (ROCHE Diagnostics, Meylan, France), prepared from a 2 $\mu\text{g ml}^{-1}$ stock solution, was added at a final concentration of 400 pg ml^{-1} . The glucocorticoid budesonide (kindly provided by Dr R. Brattsand, Astra, Lund, Sweden) was prepared from a 10 mM stock solution in absolute ethanol and used at 0.1 μM . Dexamethasone (Sigma Chemicals, St Louis, MO, U.S.A.) was used at 1 μM in similar conditions. The glucocorticoid antagonist RU486 (Sigma Chemicals) was prepared from a 10 mM stock solution in absolute ethanol and incubated at 1 μM for 1 h before IL-1 β and glucocorticoid treatments. Similar experiments were performed with staggered treatment starts, to verify the effect of the duration of the culture. Supernatants were sampled and stored at -80°C until the SCF protein assay. Fibroblasts were harvested for total RNA extraction.

For analysis of the transcript stability, confluent fibroblasts were treated with solvent alone (ethanol; 1 μM), IL-1 β (400 pg ml^{-1}), budesonide (0.1 μM) or the combination of both for 2.5 or 10 h. Supernatants were replaced by the starving medium, which contained 5 $\mu\text{g ml}^{-1}$ actinomycin D

(Sigma Chemicals). Cells were harvested at every hour from 0 to 4 h and total RNA was extracted.

SCF ELISA

Immunoreactive SCF released in the supernatant of fibroblasts treated with cytokine, glucocorticoid, and/or RU486, and the control fibroblasts treated only with solvent was quantified by a sensitive ELISA procedure. It used a capture anti-human SCF monoclonal antibody (clone 13302.6; R&D Systems Europe, Abingdon, U.K.) and a biotinylated detection anti-human SCF polyclonal antibody (R&D Systems Europe), revealed by extravidin-horseradish peroxidase (Sigma) and a 3,3',5,5'-tetramethylbenzidine liquid substrate system (Sigma Chemicals). Standard curves were generated with recombinant human SCF (R&D Systems Europe) diluted in the starving medium and were linear from 3.9 to 500 pg ml^{-1} .

Extraction of total RNA and reverse transcription

Total RNA was extracted from control or treated fibroblasts with TriReagent[®] (Molecular Research Center Inc., Cincinnati, OH, U.S.A.). Isolated RNA was diluted in RNase-free water and quantified by absorbance measurement at 260 nm. Four μg of total RNA were incubated with 0.5 μg of random primers for 5 min at 70°C and allowed to cool down to room temperature. RNA was subsequently reverse transcribed in 1 \times reverse transcription (RT) buffer (75 mM KCl, 3 mM MgCl₂, 10 mM dithiothreitol, and 50 mM Tris-HCl, pH 8.3) that contained 1 unit μl^{-1} RNasin ribonuclease inhibitor, 1 mM of each dNTP, and 10 units μl^{-1} RNase H(-)-Moloney leukaemia virus reverse transcriptase (all reagents from Promega, Madison, WI, U.S.A.). The reaction was conducted for 1 h at 37°C, and the reverse transcriptase was then heat-inactivated at 99°C for 5 min.

Quantification of SCF cDNA

Quantification of SCF cDNA was performed by polymerase chain reaction (PCR), and the results were normalized to total cDNA.

PCR of SCF cDNA SCF cDNA was amplified from total cDNA obtained after reverse transcription by online fluorescent PCR (LightCycler[®]-SYBR Green I, ROCHE Diagnostics), with primers that lead to a single 149 bp PCR product. PCR reactions were performed in 1 \times PCR reaction buffer (2 μl of the reaction mix containing FastStart Taq DNA polymerase, dNTP mix, SYBR Green I, 3 mM MgCl₂) (all from ROCHE Diagnostics), and 10 pmol of each primer: sense primer: 5'-TGGATAAGCGAGATGGTAGT-3', antisense primer: 5'-TTTTCTTTCACGCACTCCAC-3', in a 20 μl final volume for 35 cycles. Each cycle consisted of 15 s denaturation at 95°C, 10 s annealing at 53°C, and 10 s extension at 72°C. Amplified SCF cDNA was analysed on-line by fluorescence, and quantified during the exponential state.

The standard SCF cDNA was obtained from pulmonary fibroblast total cDNA. After amplification by PCR with the primers described above, the 149 bp-PCR products were electrophoresed on a 2% agarose gel stained with ethidium bromide, purified on QIAEX II (QIAEX II gel extraction kit,

QIAGEN, Courtaboeuf, France), and quantified by fluorescence (PicoGreen[®], Molecular Probes Inc.) according to a standard curve obtained with double-stranded phage λ DNA ($0-1 \mu\text{g ml}^{-1}$). This purified SCF cDNA was used as a standard curve from 1 to 300 fg ml^{-1} .

Quantification of total cDNA One μl of total cDNA was diluted (1:15) and added to $250 \mu\text{l}$ Tris-EDTA (TE) $1\times$ buffer (10 mM Tris-HCl, pH 7.8, 1 mM EDTA, pH 8). This reaction mix was incubated with $250 \mu\text{l}$ OliGreen[®] (1:200) (Molecular Probes Inc., Eugene, Oregon, U.S.A.) for 10 min in the dark. Simultaneously, a standard single-stranded M13 phage DNA (Sigma) curve ($0-500 \text{ ng ml}^{-1}$) was obtained in the same conditions. The total cDNA was quantified in a fluorometer (Hitachi F-2000, Tokyo, Japan) at an excitation wavelength of 480 nm and an emission wavelength of 520 nm. Total cDNA remained unchanged over time and for all treatments.

Expression of results and statistical analysis

SCF cDNA was calculated as $\text{fg SCF cDNA/ng total cDNA}$, and SCF protein was calculated in pg ml^{-1} . The values after treatment were expressed as a percentage of the control values. Results are the means \pm s.e.mean of $n=4$ experiments performed in duplicate. Statistical analysis of the results used Student's *t*-test, or Student-Newman-Keuls when more than two variables were compared. Values of $P<0.05$ were considered significant.

Results

Effect of IL-1 β on SCF production

SCF mRNA IL-1 β (400 pg ml^{-1}) significantly increased SCF mRNA expression by human lung fibroblasts in culture (Figure 1a,b). This increase reached +91% at 2.5 h ($128.2 \pm 51.4 \text{ fg ng}^{-1}$ total cDNA for control and 245.9 ± 99.4 for IL-1 β -treated fibroblasts; $P<0.001$, $n=4$) (Figure 1a). This effect decreased over time, but was still significant at 10 h (+39%; $75.3 \pm 23.7 \text{ fg ng}^{-1}$ total cDNA for control and 102.6 ± 36.6 for IL-1 β -treated fibroblasts; $P<0.01$, $n=4$) (Figure 1b).

SCF protein SCF protein production also increased after treatment with IL-1 β (Figure 1c,d). At 2.5 h, this increase reached +32% ($36.8 \pm 1.6 \text{ pg ml}^{-1}$ for control and 48.7 ± 3.3 for IL-1 β -treated fibroblasts; $P<0.01$, $n=4$) (Figure 1c). At 10 h, the increase was less but still significant (+16%; $74.3 \pm 12.1 \text{ pg ml}^{-1}$ for control and 85.7 ± 14.6 for IL-1 β -treated fibroblasts; $P<0.05$, $n=4$) (Figure 1d). Similar results were obtained when IL-1 β treatment was staggered to stop the cell culture at the same time. This finding confirmed that the IL-1 β effect we observed was not due to a modification in the cell phenotype over time.

Effect of budesonide on IL-1 β -induced SCF production

SCF mRNA Budesonide ($0.1 \mu\text{M}$) inhibited the IL-1 β -induced increase in SCF mRNA expression (Figure 1a,b). This inhibition reached 68% at 2.5 h ($165.7 \pm 64.1 \text{ fg ng}^{-1}$

total cDNA; $P<0.01$, $n=4$) (Figure 1a) and continued so that this expression was below the baseline at 10 h (192% inhibition; $42.4 \pm 4.4 \text{ fg ng}^{-1}$ total cDNA; $P<0.01$, $n=4$) (Figure 1b). Conversely, as previously shown (Kassel *et al.*, 1998), budesonide alone caused SCF mRNA expression to decrease below baseline at 2.5 h (-36%; $85.7 \pm 35.7 \text{ fg ng}^{-1}$ total cDNA; $P<0.001$, $n=4$) (Figure 1a), whereas it provoked a slight increase at 10 h (+17% compared with baseline expression; $91.5 \pm 30.1 \text{ fg ng}^{-1}$ total cDNA; $P<0.05$, $n=4$) (Figure 1b).

SCF protein The modulation by budesonide over time was different for the IL-1 β -induced increase of SCF protein than for that of SCF mRNA (Figure 1c,d). A surprising increase over IL-1 β -induced expression alone was observed at 2.5 h (61%; $56.0 \pm 1.0 \text{ pg ml}^{-1}$ SCF; $P<0.001$, $n=4$) (Figure 1c). As was expected from the mRNA modulation, however, total inhibition of the effect of IL-1 β was observed at 10 h (-9% as compared with baseline expression; $64.6 \pm 9.3 \text{ pg ml}^{-1}$ SCF; $P<0.05$, $n=4$) (Figure 1d). Conversely, budesonide alone caused SCF protein expression to decrease at 2.5 h (-13%; $32.1 \pm 1.7 \text{ pg ml}^{-1}$ SCF; $P<0.01$, $n=4$) (Figure 1c) and at 10 h (-11%; $66.0 \pm 13.4 \text{ pg ml}^{-1}$ SCF; $P<0.05$, $n=4$) (Figure 1d).

Results were similar with dexamethasone (Table 1): the effect of the glucocorticoids was totally opposite in constitutive and inflammatory conditions.

Effect of RU486 on SCF expression

To verify that these effects depended on the glucocorticoid receptor, we evaluated the effect of the glucocorticoid-receptor antagonist RU486 on the budesonide-induced effects.

SCF mRNA (Figure 1a,b) RU486 ($1 \mu\text{M}$) by itself had no effect on SCF mRNA constitutive expression ($130.3 \pm 53.0 \text{ fg ng}^{-1}$ total cDNA; $n=4$ at 2.5 h; $79.0 \pm 23.1 \text{ fg ng}^{-1}$ total cDNA, $n=4$ at 10 h). In addition, RU486 antagonized both effects of budesonide: the decrease in SCF mRNA expression at 2.5 h ($138.2 \pm 56.7 \text{ fg ng}^{-1}$ total cDNA for RU486 + budesonide-treated fibroblasts; $P<0.05$, $n=4$) (Figure 1a), and the increase induced by budesonide alone at 10 h ($75.1 \pm 21.1 \text{ fg ng}^{-1}$ total cDNA; $n=4$) (Figure 1b), as well as the inhibiting effect of budesonide on the IL-1 β -stimulated SCF mRNA expression at 2.5 h ($197.7 \pm 3.8 \text{ fg ng}^{-1}$ total cDNA; NS, $n=4$) (Figure 1a) and at 10 h ($91.3 \pm 22.1 \text{ fg ng}^{-1}$ total cDNA; NS, $n=4$) (Figure 1b). However, RU486 alone slightly decreased the IL-1 β -stimulated SCF mRNA expression at 2.5 h by 28% ($212.6 \pm 83.6 \text{ fg ng}^{-1}$ total cDNA for RU486 + IL-1 β -treated fibroblasts; $P<0.001$, $n=4$) (Figure 1a), but not at 10 h ($92.6 \pm 23.0 \text{ fg ng}^{-1}$ total cDNA; NS, $n=4$) (Figure 1b).

SCF protein expression (Figure 1c,d) RU486 had no effect on SCF constitutive protein expression ($35.0 \pm 2.2 \text{ pg ml}^{-1}$ SCF; $n=4$ at 2.5 h; $74.6 \pm 12.1 \text{ pg ml}^{-1}$ SCF; $n=4$ at 10 h). Similarly, it totally cancelled out the effect of budesonide alone at 2.5 h ($37.4 \pm 1.5 \text{ pg ml}^{-1}$ SCF; $n=4$) (Figure 1c) and at 10 h ($75.7 \pm 11.0 \text{ pg ml}^{-1}$ SCF; $n=4$) (Figure 1d). It also cancelled out the effect of budesonide on the IL-1 β -enhanced SCF protein expression at 2.5 h ($48.0 \pm 0.3 \text{ pg ml}^{-1}$ SCF; NS, $n=4$) (Figure 1c) and the inhibition induced at 10 h

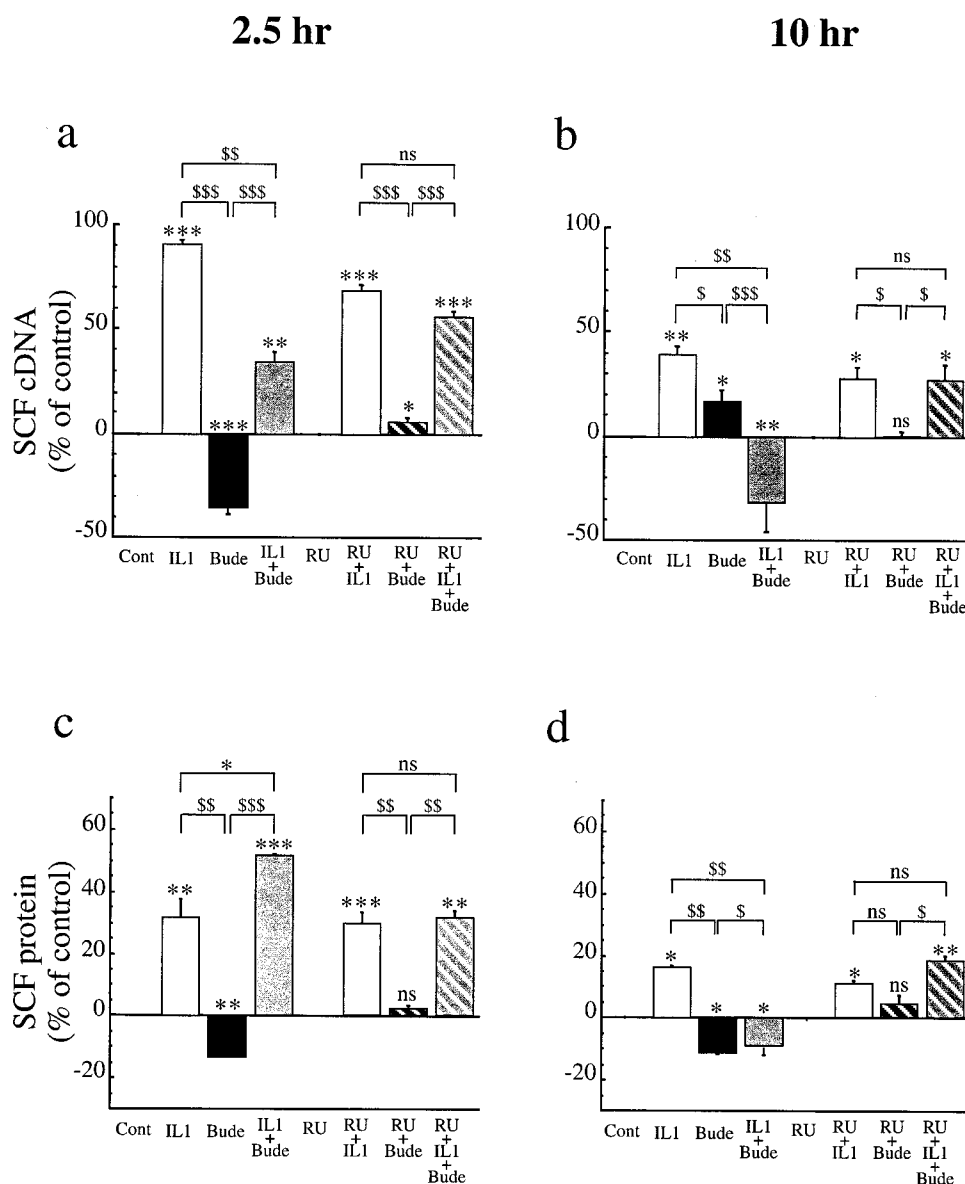


Figure 1 Effect of IL-1 β (IL1, 400 pg ml⁻¹; pale grey) or budesonide (bude, 0.1 μ M; black) or IL-1 β + budesonide (IL1 + bude; dark grey) and/or RU486 (1 μ M; hatched columns) on SCF mRNA (a,b) and protein (c,d) expression at 2.5 h (a,c) and 10 h (b,d) by human lung fibroblasts in culture. SCF cDNA was quantified after total RNA reverse transcription by online fluorescent PCR. Results were normalized to total cDNA and expressed as fg SCFcDNA/ng total cDNA; the values after treatment were expressed as a percentage of the control values. SCF protein levels were assessed in the supernatant by ELISA and the values after treatment expressed as a percentage of the control values, in pg ml⁻¹. Results are mean values (blocks) \pm s.e.mean (bars) of four experiments performed in duplicate on fibroblasts from three different donors (* P < 0.05; ** P < 0.01; *** P < 0.001).

(87.8 \pm 0.7 pg ml⁻¹ SCF; NS, n = 4) (Figure 1d). Finally, RU486 had no effect on the IL-1 β -enhanced SCF protein expression at 2.5 h (48.0 \pm 3.0 pg ml⁻¹ SCF; P < 0.001, n = 4) (Figure 1c) or at 10 h (80.3 \pm 14.1 pg ml⁻¹ SCF; P < 0.05, n = 4) (Figure 1d). Similar results were observed with dexamethasone (Table 1).

SCF mRNA stability

The mechanism of the glucocorticoid-induced effects in IL-1 β -treated fibroblasts was analysed by evaluating the stability of the SCF mRNA by measuring its half-life ($t_{1/2}$) after adding actinomycin D.

At 2.5 h (Figure 2a), the estimated half-life of SCF mRNA was 2.0 h in the control fibroblasts treated with solvent alone. IL-1 β treatment significantly increased SCF mRNA stability to $t_{1/2}$ = 3.8 h (compared with control, P < 0.001). The addition of budesonide slightly decreased this enhanced stability, to $t_{1/2}$ = 3.1 h, still significantly higher than the control (P < 0.001). Budesonide alone, on the other hand, significantly decreased the stability of the SCF mRNA to $t_{1/2}$ = 1.2 h (compared with control, P < 0.001).

At 10 h (Figure 2b), the estimated half-life was 3.4 h in the control cells, thereby showing that SCF mRNA stability increases with time. Treatment with either IL-1 β or budesonide did not significantly modify this stability

Table 1 Effect of dexamethasone on SCF mRNA and protein expression

SCF mRNA ($fg\ ng^{-1}$)	at 2.5 h		at 10 h	
	-RU486	+RU486	-RU486	+RU486
Control	103.8 ± 18.6	106.6 ± 18.1	136.4 ± 12.5	138.0 ± 9.3
IL-1 β	163.8 ± 18.9 <i>P</i> < 0.01	147.9 ± 11.0 <i>P</i> < 0.01	189.9 ± 16.7 <i>P</i> < 0.001	160.1 ± 15.6 <i>P</i> < 0.001
Dexamethasone	83.1 ± 13.6 <i>P</i> < 0.01	106.8 ± 12.7 ns	137.9 ± 26.3 ns	133.4 ± 19.4 ns
IL-1 β + dexamethasone	107.7 ± 24.8 ns	141.8 ± 4.1 <i>P</i> < 0.05	69.1 ± 7.2 <i>P</i> < 0.001	158.9 ± 10.5 <i>P</i> < 0.001
SCF protein ($pg\ ml^{-1}$)	-RU486	+RU486	-RU486	+RU486
Control	32.4 ± 3.0	34.0 ± 2.1	65.7 ± 7.3	62.2 ± 3.2
IL-1 β	43.2 ± 2.3 <i>P</i> < 0.01	40.6 ± 2.1 <i>P</i> < 0.01	80.1 ± 9.0 <i>P</i> < 0.05	78.1 ± 8.1 <i>P</i> < 0.01
Dexamethasone	29.0 ± 3.0 <i>P</i> < 0.05	32.6 ± 3.0 ns	60.1 ± 4.2 ns	70.1 ± 8.5 ns
IL-1 β + dexamethasone	45.7 ± 7.0 <i>P</i> < 0.05	40.3 ± 1.5 <i>P</i> < 0.01	52.5 ± 3.3 <i>P</i> < 0.05	81.3 ± 0.3 <i>P</i> < 0.01

Effect of IL-1 β (400 pg ml⁻¹), dexamethasone (1 μ M) and/or RU486 (1 μ M) on SCF mRNA and protein expression by human lung fibroblasts in culture at 2.5 h and 10 h. SCF cDNA was quantified by on-line fluorescent PCR, and SCF protein by ELISA. Results were normalized to total cDNA, which did not vary over time. Results are expressed as (SCF cDNA)/(total cDNA) in fg ng⁻¹. SCF protein levels were assessed in the supernatant by ELISA and results are expressed as (SCF protein) in pg ml⁻¹. Results are mean values \pm s.e. mean of four experiments performed in duplicate on fibroblasts from three different donors.

SCF mRNA stability

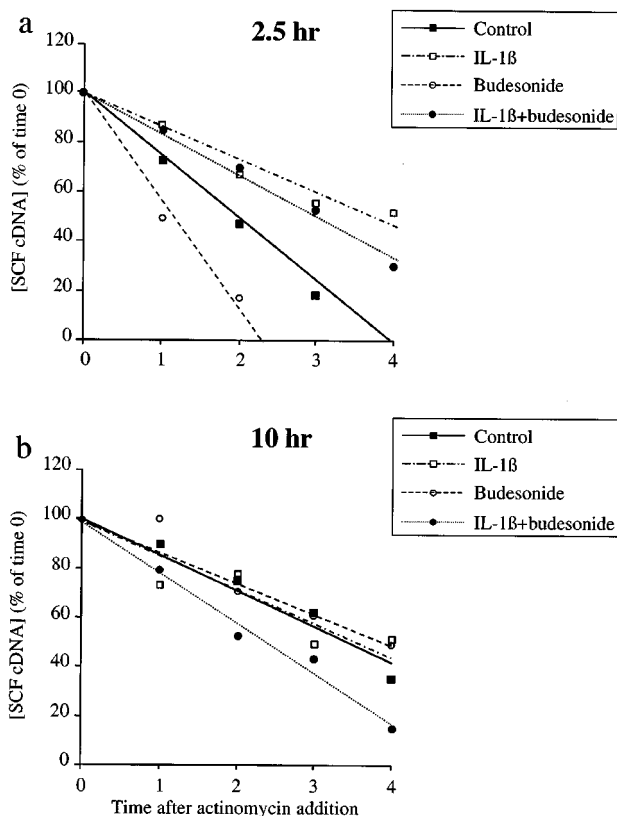


Figure 2 SCF mRNA stability. Fibroblasts were treated with IL-1 β (400 pg ml⁻¹), budesonide (0.1 μ M), IL-1 β plus budesonide or ethanol (1 μ M; control) for 2.5 h (a) or 10 h (b). Fibroblasts were then washed and treated with 5 μ g ml⁻¹ actinomycin D for the indicated time. SCF cDNA was quantified by on-line fluorescent PCR. Results are expressed as a percentage of the quantity of SCF cDNA measured for each treatment at time 0 of actinomycin D addition, which is represented as the 100% of each treatment. Data are means of three experiments performed in duplicate on fibroblasts from three different donors.

($t_{1/2}$ = 3.5 h and 3.8 h, respectively, NS). The combination of IL-1 β and budesonide markedly decreased it, however ($t_{1/2}$ = 2.4 h, *P* < 0.001).

Discussion

We present here evidence that (1) the pro-inflammatory cytokine IL-1 β substantially increases the expression of SCF mRNA and protein by human lung fibroblasts in culture, and (2) that this increased expression is markedly regulated by the glucocorticoids budesonide and dexamethasone.

This first finding, that IL-1 β , present in large amounts in the human airways of asthmatic patients (Borish *et al.*, 1992), heightens SCF mRNA expression, is in agreement with the reported increase in the IL-1 β -mediated expression of genes implicated in the inflammatory process, such as cyclooxygenase (COX)-2 mRNA expression by human pulmonary epithelial cells (Lin *et al.*, 2000) or eotaxin mRNA expression in A549 airway epithelial cells (Jedrzkiewicz *et al.*, 2000). Both reports propose a mechanism for this increased expression associated with the activation of the transcription factor nuclear factor (NF)- κ B by IL-1 β (Jedrzkiewicz *et al.*, 2000; Lin *et al.*, 2000); this mechanism would involve gene promoters with at least one NF- κ B responsive element. One such element has been identified in the SCF gene promoter (Taylor *et al.*, 1996), although its specific role in SCF gene expression is not yet known.

We also observed that when SCF expression increased, its mRNA became more stable (had a longer half-life). This is consistent with the increased stability of COX-2 mRNA observed in HeLa cells stimulated by IL-1 β (Ridley *et al.*, 1998) and of inducible nitric oxide synthase mRNA in cardiac myocytes (Oddis *et al.*, 1995). Among the mechanisms involved in increased mRNA stability is the length of the poly(A)⁺ tail (Ross, 1995), and it has been proposed that IL-1 β increases the process of polyadenylation (Stoeckle, 1992), which might play a part in this effect. Accordingly, SCF

mRNA and protein behave like many other proinflammatory factor genes: their expression increases in the presence of IL-1 β . These genes include COX-2 (Lin *et al.*, 2000), RANTES (Saji *et al.*, 2000), IL-6 and IL-8 (Agarwal *et al.*, 1995), eotaxin (Ghaffar *et al.*, 1999) and NGF (Olgart & Frossard, 2001).

Budesonide is the major anti-inflammatory compound used in the treatment of diseases such as asthma that are associated with an increased number of tissue mast cells. Here we studied its effect on SCF expression in inflammatory conditions created by IL-1 β , rather than the previously-reported effects on its constitutive expression (Kassel *et al.*, 1998). We showed here that budesonide inhibited the IL-1 β -induced increase in SCF mRNA and protein expression in the short term (2.5 h) and even more so longer after treatment (10 h). This effect is consistent with many others reported in the literature, where glucocorticoids inhibit the expression of genes involved in the inflammatory process, including IL-4 and 5 (Bentley *et al.*, 1996) or IL-1 β (Sousa *et al.*, 1997) (for review, see Barnes *et al.*, 1998). We have demonstrated here that this inhibitory effect involves the destabilization of SCF mRNA in the short term and its aggravation after a longer period.

The precise mechanisms of glucocorticoids action in SCF mRNA stability is unclear. It has been suggested that AU-rich elements in the 3'-untranslated region play some role in the glucocorticoid-induced decrease in mRNA stability (Peppel *et al.*, 1991; Ristimaki *et al.*, 1996), but none have been found in the human SCF mRNA sequence (Martin *et al.*, 1990). Hence, the observed SCF mRNA destabilization probably involves another mechanism, as yet unidentified. Possibilities include the down-regulation of Poly(A)⁺ binding proteins (PABP) or dysregulation of the 5'-Catabolite gene Activator Protein (CAP) that protects mRNA from degradation (Ross, 1995). Other currently unknown mechanisms may also play a role in the inhibition by budesonide of IL-1 β -stimulated SCF expression. Since the SCF gene promoter has NF- κ B and AP-1 responsive elements (Taylor *et al.*, 1996), a direct interaction between activated GR and these transcription factors may be involved. This interaction has been proposed as a general mechanism (Adcock, 2001) and been shown to exist in specific cases between GR and NF- κ B, involving repression of histone acetyl transferase activity activated by p65 (Ito *et al.*, 2001) and between GR and AP-1 (Diamond *et al.*, 1990; Kerppola *et al.*,

1993). Nevertheless, we also noticed that the short-term increase in SCF protein expression induced by the combination of IL-1 β and budesonide was substantially greater than that induced by IL-1 β alone, conversely to the decreased SCF mRNA expression. This surprising effect may be related to an earlier increase in SCF mRNA expression, to a rapid increase of the translational process after short term treatment by glucocorticoids, or to both. However, SCF gene expression is not described in the literature as an early gene and thus not regulated in the same way as the early gene family. The mechanism of this interesting event requires further investigation.

Additionally, we have demonstrated here that all the results obtained with budesonide were glucocorticoid-receptor dependent, since RU486, a glucocorticoid-receptor antagonist (Cairns *et al.*, 1993), cancelled out all the effects induced by budesonide on SCF mRNA and protein expression, whether or not IL-1 β was present. RU486 alone exerted a slight inhibitory effect on IL-1 β -induced SCF mRNA expression. Since RU486 had no effect on control cells or on SCF mRNA stability (not shown), we may hypothesize that upon IL-1 β stimulation, RU486 interacts with the transductional pathways activated by IL-1 β . In accordance with this hypothesis, it has been reported that RU486, after binding to the glucocorticoid receptor (GR), interacts with the transcription factor NF- κ B and inhibits NF- κ B activity without affecting the latter's binding to DNA (Hofmann *et al.*, 1998; Liden *et al.*, 1997), through interaction with a histone acetyltransferase/NF- κ B complex (Ito *et al.*, 2001). A similar mechanism may thus be proposed to explain the decrease in IL-1 β -enhanced SCF mRNA expression shown in this study.

In conclusion, our study clearly shows that IL-1 β , a pro-inflammatory cytokine present in large quantities in the airways of asthmatic patients, and here mimicking the inflammatory conditions in human lung fibroblasts *in vitro*, increases SCF expression, at least partly by increasing the stability of its mRNA. Moreover, anti-inflammatory glucocorticoids decreased this IL-1 β -enhanced SCF expression and the stability of its mRNA. This may be of major importance in the mechanism of glucocorticoid treatment of diseases associated with an increased number of mast cells in tissues, such as asthma. Our results thus suggest a new mechanism of glucocorticoid action, through the regulation of the expression of the mast cell growth factor SCF.

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