

Modulation of ICAM-1 expression in human alveolar macrophages *in vitro*

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Modulation of ICAM-1 expression in human alveolar macrophages in vitro. M. Fattal-German, F. Le Roy Ladurie, J. Cerrina, F. Lecerf, S. Berrih-Aknin. ©ERS Journals Ltd 1996.

ABSTRACT: Modulation of intercellular adhesion molecule-1 (ICAM-1) expression may be a basic mechanism by which alveolar macrophages (AMs) regulate the inflammatory process in the lung in response to local stimuli.

As a model for studying the anti-inflammatory activity of drugs on human AMs, we investigated the effects of fusafungine, an antibiotic for local use by aerosol with anti-inflammatory properties, and that of the glucocorticoid dexamethasone, on ICAM-1 expression induced *in vitro* by recombinant interferon- γ (rIFN- γ). ICAM-1 protein expression was studied on AMs by means of flow cytometry with an anti-CD54 monoclonal antibody; messenger ribonucleic acid (mRNA) levels were determined by reverse transcriptase-polymerase chain reaction (RT-PCR).

ICAM-1 was expressed before culture on 21% of bronchoalveolar lavage (BAL) cells, with low intensity. Culture for 24 h with rIFN- γ resulted in a significant increase in ICAM-1 protein expression (82% of cells were strongly positive). Fusafungine significantly inhibited rIFN- γ -induced ICAM-1-protein expression on AMs in a concentration-dependent fashion. The mechanism of ICAM-1 downregulation was mainly post-transcriptional, but also partly transcriptional. By contrast, dexamethasone did not influence rIFN- γ -induced ICAM-1 expression.

This *in vitro* model using human AMs should prove useful for investigating the cellular and molecular targets of anti-inflammatory drugs.

Eur Respir J., 1996, 9, 463–471.

Macrophages are resident tissue cells that play a major role in natural and acquired immunity. Alveolar macrophages (AMs), in addition to many other important functions, participate in the modulation of immune responses as antigen-presenting cells (APCs). They thus possess surface molecules that enable them to recognize and adhere to lymphocytes. In addition to presentation of antigen to the T-cell receptor, several antigen-nonspecific accessory molecules promote attachment of T-cells to APC; this is an important step in the induction of the immune response, through the transmission of regulatory signals to the T-cell, and the induction of target cell functional activity [1].

Intercellular adhesion molecules (ICAMs) on the APC surface, together with leucocyte function-associated molecule-1 (LFA-1) on T-cells, form a critical adhesive ligand-receptor pair. Interaction of LFA-1 and ICAMs is required to activate resting T-cells, and to induce activated T-cells to proliferate and function optimally. The ICAM class currently contains three molecules. The complementary deoxyribonucleic acids (cDNAs) for the three ICAMs have been cloned and the corresponding amino acid sequences have been determined [2–4]. There is a high degree of sequence homology between the extracellular portions (*e.g.* 48% between ICAM-3 and ICAM-1 [5]). However, these molecules differ in their

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Keywords: Alveolar macrophages
anti-inflammatory drugs
intercellular adhesion molecule-1

Received: May 18 1995
Accepted after revision November 27 1995

This work was supported by grants from AFLM, Université Paris-Sud and IRIS.

tissue distribution, binding activity for LFA-1, and the functions of the intracytoplasmic region. ICAM-1 has been reported to bind LFA-1 with an avidity several times greater than the other ICAMs [6]. The main functional role of ICAM-1 and ICAM-2 is merely to bind to LFA-1, the only molecule transducing signals that enhance cell activation. Only ICAM-3 has two clear functions, as it binds the LFA-1 molecule and is also able to transduce signals. Interestingly, ICAM-1 (CD54) is expressed weakly on normal resting AMs. However, it can be induced on monocytes/macrophages in culture following activation with inflammatory mediators, including lipo-polysaccharide (LPS), 12-O-tetradecanoyl phorbol-13-acetate (TPA), interleukin-1 (IL-1), interleukin-3 (IL-3), interleukin-4 (IL-4), recombinant human granulocyte/macrophage colony-stimulating factor (GM-CSF), tumour necrosis factor- α (TNF- α) and recombinant gamma-interferon (rIFN- γ) [7–12].

Regulation of ICAM-1 expression may be one of the basic mechanisms by which AMs modulate immune and inflammatory processes in the lung in response to local stimuli, through interactions between AMs and T-cells [13]. As a model to study the anti-inflammatory activity of drugs on AMs, we investigated the effects of fusafungine, an antibiotic for local use by aerosol with anti-inflammatory properties [14–17], and that of the glucocorticoid,

dexamethasone, on rIFN- γ -induced ICAM-1 expression in cultured human AMs. Expression was studied in terms of the surface antigen (immunocytofluorimetry) and the messenger ribonucleic acid (mRNA) (reverse transcriptase-polymerase chain reaction (RT-PCR)).

Materials and methods

Reagents

AMs were cultured in RPMI 1640 medium (Gibco-BRL) containing 2 mM L-glutamine, 25 mM hydroxyethylpiperazine ethanesulphonic acid (HEPES) (Gibco-BRL), 100 U·mL⁻¹ penicillin G, 100 μ g·mL⁻¹ streptomycin, 5 μ g·mL⁻¹ fungizone, 50 μ g·mL⁻¹ amikacin, and 10% human AB serum.

rIFN- γ was a gift from Roussel-Uclaf (Paris). Stock rIFN- γ was prepared at 1 \times 10⁶ U·mL⁻¹ in sterile supplemented RPMI-1640.

Fusafungine (IRIS, France) was solubilized in dimethyl sulphoxide (DMSO) at 10 mg·mL⁻¹, and twofold dilutions were made in supplemented RPMI medium with vigorous shaking to obtain solutions ranging from 320 to 10 μ g·mL⁻¹, *i.e.* 20 times the final concentrations in the assay.

Dexamethasone (Sigma) was stored as a stock solution of 5 \times 10⁻³ M in ethanol at -20°C.

Bronchoalveolar lavage (BAL)

BAL for diagnosis or routine monitoring was performed in the subsegmental bronchus of the middle lobe of patients treated at Hôpital Marie-Lannelongue for lung diseases. With the patient's informed consent, three 50 mL aliquots of sterile 0.9% saline were instilled, aspirated manually and pooled. Samples remaining after cytological, bacteriological and virological tests were used in this study. Samples were kept on ice, and then filtered through a single layer of cotton gauze. Total and differential cell counts were performed. Cell viability always exceeded 85% (trypan blue exclusion). BAL samples containing bacteria (direct examination) or more than 50% neutrophils were discarded.

Macrophage culture

BAL fluid was centrifuged at 300 \times g for 10 min at 4°C, and cells were resuspended in supplemented RPMI. AMs were adherence purified, and then 1 \times 10⁶ AMs were incubated at 37°C with 5% CO₂ in 24-well plates with rIFN- γ (250 U·mL⁻¹). The effects of rIFN- γ and that of the drugs on AM activation measured by ICAM-1 expression were investigated after 24 h of incubation. For this, nonadherent cells were discarded and, after gentle washing with medium, adherent AMs were detached from the plates using cold phosphate-buffered saline (PBS) containing 0.02% ethylenediamine tetra-acetic acid (EDTA) and vigorous pipetting.

Flow cytometric phenotype analysis

ICAM-1 expression on the surface of AMs was investigated before and after culture, by means of flow cytometry with indirect immunofluorescence [18]. The specific monoclonal antibodies (MoAbs) used were as follows: IOT2a (mouse-immunoglobulin G_{2a} (IgG_{2a}), conjugated with fluorescein isothiocyanate (FITC); Immunotech), which binds a common nonpolymorphic human leucocyte antigen-DR (HLA-DR) epitope; and IOL54 (unconjugated mouse-IgG₁; Immunotech), that reacts with the ICAM-1 antigen (CD54). Fluorescence data were collected in a gate of 5,000 cells selected according to size and granularity, using a fluorescence-activated cell sorter (FACSscan) apparatus (Becton Dickinson) with an argon laser operating at 488 nm. In this gate, the proportion of HLA-DR-positive cells ranged 75–98% (mean (SD) 87 (9)%). Data were analysed using Lysis-2 software (Becton Dickinson). Fluorescence associated with anti-CD54 MoAbs on activated cells clearly showed two peaks, one corresponding to weakly positive cells and the second to strongly positive cells. ICAM-1 fluorescence is expressed as the percentage of strongly positive cells (fluorescence intensity >30 on the logarithmic scale). Mean fluorescence intensity (mFI) was also used as an index of ICAM-1 density on the cell surface. Percentage inhibition was calculated as follows: (% CD54+ cells after culture without fusafungine - % CD54+ cells after culture with fusafungine) \times 100/(% CD54+ cells after culture without fusafungine - % CD54+ cells before culture).

ICAM-1 mRNA detection

ICAM-1 mRNA was assayed by means of polymerase chain reaction (PCR) on complementary deoxyribonucleic acid (cDNA) derived enzymatically from ribonucleic acid (RNA) extracted from 1 \times 10⁶ fresh or cultured BAL cells. Cells were pelleted by centrifugation and resuspended in 400 μ L of RNazol (Bioprobe, Paris), which lyses cells and protects RNA from degradation, and were kept at -80°C until RNA extraction.

Total RNA extraction. Total RNA was extracted from AMs by a guanidinium thiocyanate-phenol-chloroform method [19].

Primers. All primers were synthesized by Genset (Paris). For ICAM-1 PCR a pair of 20 nucleotide primers, located in the 150–348 nucleotide sequence of the I domain of ICAM-1 cDNA, were chosen to amplify a 189 base pair (bp) fragment. This sequence has no homology with NCAM-1 cDNA, and very little homology with ICAM-2 or ICAM-3 cDNAs. The primer sequences were 5' AAA GTC ATC CTG CCC CGG GG 3' (sense) and 5' AGG GCA GTT TGA ATA GCA CA 3' (antisense). The latter was used for specific reverse transcription. The fusion temperatures were 66 and 62°C, respectively.

To check for the presence of cDNA in all the negative samples and for semiquantitative assay of ICAM-1 mRNA, glyceraldehyde-3-phosphate dehydrogenase (GAPDH) mRNA was investigated in parallel, using the

following primers: 5' ATC ACC ATC TTC CAG GAG CG 3' (sense), and 5' CCT GCT TCA CCA CCT TCT TG 3' (antisense). This latter primer was used for specific reverse transcription of mRNA into GAPDH cDNA. This pair of 20 nucleotide primers amplifies a 573 bp fragment.

Specific reverse transcription. To convert mRNA into cDNA for PCR, total RNA (1 µg) was first heated to 72°C for 10 min and cooled on ice. The mRNA was then reverse transcribed in a 50 µL reaction volume containing buffer (Gibco-BRL), 10 mM dithiothreitol, 20 U of ribonuclease (RNase) inhibitor (RNAsin, Boehringer), 0.5 mM (each) deoxynucleotide triphosphate (dNTP) (Bioprobe) mix, 40 µM specific primer, and 50 U of Molony murine leukaemia virus (MMLV) reverse transcriptase (Gibco-BRL), for 1 h at 37°C in a water bath. The enzyme was then denatured by incubation at 95°C for 5 min.

PCR for cDNA amplification. cDNA (5 µL) was amplified in a 50 µL reaction containing buffer (20 mM Tris-HCl, pH 8.55, 16 mM (NH₄)₂SO₄, 150 µg·mL⁻¹ bovine serum albumin (BSA), 1.25 mM MgCl₂) containing 0.2 mM each dNTP, 0.5 µM each primer, and 1.25 U of TAQ DNA-polymerase (Eurobio, Les Ulis, France). The samples were amplified in a thermocycler (PHC-3, Techne) with the following parameters: 94°C, 1 min; 63°C, 1 min; 72°C, 1 min; 40 cycles). A final extension step was performed at 72°C for 1.5 min. Positive and negative controls were analysed in all experiments. mRNA extracts not submitted to reverse transcription were also included to check the absence of contaminating DNA. All samples were analysed at least three times. GAPDH mRNA was assessed in the same conditions.

Analysis of PCR products. PCR products (10 µL) were analysed by electrophoresis on 2% agarose gel containing 0.5 µg·mL⁻¹ ethidium bromide. The bp size marker was the pUC 18 Hae III digest (Sigma). GELS were photographed under ultra violet (UV) light. For semiquantitative assessment of band intensity, the gels were photographed with

a video imager (Imstar, Paris) in reverse mode, stained DNA appearing as black bands. The relative amount of DNA was determined densitometrically by using Starwise Akigel software (Imstar, Paris). Results for each sample are expressed as arbitrary optical density (OD) units related to the integrated volume calculated on the densitometer.

Statistical analysis

Comparisons were made using Student's two-sided t-test. Linear regression analysis was used to identify correlations. Differences with p-values lower than 0.05 were considered significant.

Results

A total of 14 BAL samples were studied. Of these, two samples were excluded because AMs were highly activated, as more than 70% of AMs (74 and 77%, respectively) expressed the ICAM-1 antigen before culture, although at a low level. Cell counts ranged 95–30% (mean 74±21%), 35–2% (mean 10±10%) and 60–1% (mean 16±19%) for AMs, neutrophils and lymphocytes, respectively.

Basal ICAM-1 expression on alveolar macrophages

The ICAM-1 molecule was expressed on 5–42% of gated cells (mean (SD) 21(13)%). ICAM-1 antigen density on the cell surface expressed as the mFI ranged 35–137 (overall mean (SD) of mFIs 66.7 (29.3)).

Fusafungine inhibits rIFN γ - induced ICAM-1 protein expression on AMs in vitro

Cells from 10 BAL specimens were incubated for 24 h with rIFN- γ (250 U·mL⁻¹) and various concentrations of fusafungine (0.5–16 µg·mL⁻¹). Cell viability always exceeded 85% (trypan blue exclusion) and cell size was unaffected by fusafungine. Flow cytometry showed that

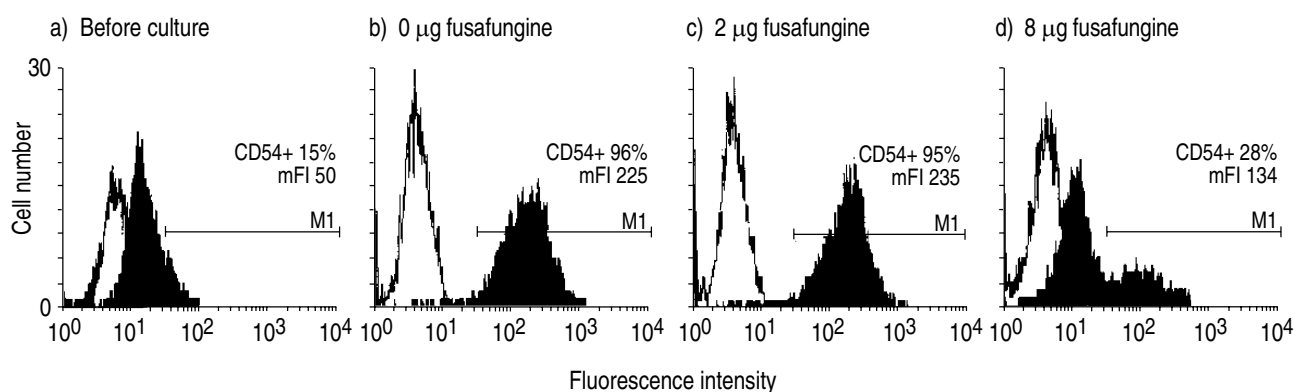


Fig. 1. – Effect of fusafungine on ICAM-1 protein expression by alveolar macrophages stimulated by rIFN- γ from one representative BAL sample. Flow cytometric analysis of ICAM-1-specific immunofluorescence with an anti-CD54 MoAb (right histogram) compared with control fluorescence (left histogram). Results are expressed as the percentage of CD54-positive cells and the mean fluorescence intensity: a) before culture; b) after 24 h of culture with 250 µ·mL⁻¹ rIFN- γ ; c) after 24 h of culture with 250 U·mL⁻¹ rIFN- γ and 2 µg·mL⁻¹ of fusafungine; d) after 24 h of culture with 250 U·mL⁻¹ rIFN- γ and 8 µg·mL⁻¹ of fusafungine. ICAM-1: intercellular adhesion molecule-1; BAL: bronchoalveolar lavage; MoAb: monoclonal antibody; rIFN- γ : recombinant interferon- γ ; mFI: mean fluorescence intensity.

ICAM-1 expression on AMs was markedly increased after 24 h of culture with rIFN- γ , since it was observed on $82\pm 16\%$ of cells ($p<0.001$ vs values before culture), with a high intensity (mFI= 172.6 ± 74.5 ; $p<0.001$ vs values before culture). ICAM-1 was expressed on 87–100% of HLA-DR-positive cells. The percentage of ICAM-1-positive cells fell in a fusafungine concentration-dependent fashion (fig. 1). Fusafungine inhibition of ICAM-1 expression on BAL cells (see Materials and Methods) was detectable at $2\ \mu\text{g}\cdot\text{mL}^{-1}$ (mean % inhibition 28 (SD 32)% inhibition; $p<0.005$ vs $0\ \mu\text{g}\cdot\text{mL}^{-1}$ fusafungine), and

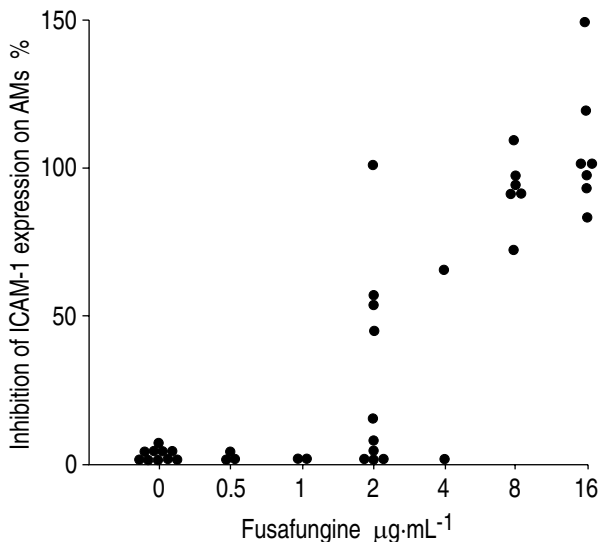


Fig. 2. – Inhibitory effect of fusafungine on ICAM-1 protein expression by alveolar macrophages from 10 BAL samples after 24 h of culture with $250\ \text{U}\cdot\text{mL}^{-1}$ rIFN- γ . Each point represents one determination, all 10 BAL were tested at $0\ \mu\text{g}\cdot\text{mL}^{-1}$. For abbreviations see legend to figure 1.

increased with concentration (fig. 2). Of note, because of the small number of AMs available due to the poor volume of BAL remaining after clinical investigations, every drug concentration could not be tested on all 10 BAL. Thus, in figure 2, all 10 BAL were tested only for the control without drug. The mean inhibition was 92% (SD 11) at $8\ \mu\text{g}\cdot\text{mL}^{-1}$ ($p<0.001$ vs $0\ \mu\text{g}\cdot\text{mL}^{-1}$) and inhibition was total at $16\ \mu\text{g}\cdot\text{mL}^{-1}$ ($p<0.001$). As illustrated in figure 1, fusafungine also reduced the mFI in parallel to the percentage of positive cells.

It is noteworthy that AMs from BAL samples Nos 7 and 8 were inhibited by more than 100% (133 and 120% inhibition, respectively) by $16\ \mu\text{g}\cdot\text{mL}^{-1}$ fusafungine. Indeed, ICAM-1 was expressed by a large proportion of these AMs before culture (40 and 42%, respectively) (table 1) but at a low intensity (mFI = 44 and 35, respectively); after 24 h culture with rIFN- γ and $16\ \mu\text{g}\cdot\text{mL}^{-1}$ of fusafungine, values were lower than before culture (29 and 19%, respectively).

ICAM-1 protein expression induced by culture with rIFN- γ was, thus, blocked by fusafungine, which also appeared to further downregulate basal ICAM-1 expression on activated AMs.

Fusafungine inhibits rIFN- γ -induced ICAM-1 mRNA expression in AMs in vitro

To determine whether fusafungine modulates the induction of ICAM-1 expression at the transcriptional or post-transcriptional level, ICAM-1 mRNA was tested for in AMs from 6 BAL samples before and after 24 h of culture, by applying PCR to ICAM-1 cDNA obtained after reverse transcription of mRNA. As shown in table 1, ICAM-1 mRNA transcripts were present before cul-

Table 1. – Effect of fusafungine on the expression of ICAM-1 antigen and mRNA in AMs from BAL (n=6)

BAL No.	ICAM-1 expression	Before culture	After 24 h culture with fusafungine $\mu\text{g}\cdot\text{mL}^{-1}$			
			0	2	8	16
5	CD54+ cells %	5	39	37	7	ND
	mFI	56	350	360	162	ND
	ICAM-1 mRNA*	0.10	0.3	0.11	0.10	ND
6	CD54+ cells %	22	85	75	27	ND
	mFI	81	195	182	160	ND
	ICAM-1 mRNA*	0.21	0.92	0.31	0.23	ND
7	CD54+ cells %	40	100	68	ND	29
	mFI	44	88	59	ND	39
	ICAM-1 mRNA*	0.21	0.69	0.54	ND	0.24
8	CD54+ cells %	42	85	40	ND	19
	mFI	35	89	111	ND	84
	ICAM-1 mRNA*	0.37	1.07	1.08	ND	0.87
9	CD54+ cells %	34	94	94	28	39
	mFI	43	231	235	140	146
	ICAM-1 mRNA*	0	0.29	0.48	0.37	0.12
10	CD54+ cells %	9	73	71	ND	10
	mFI	42	208	235	ND	210
	ICAM-1 mRNA*	0.14	0.72	0.98	ND	0.16

*: expressed as the intensity of optical density (OD) of the 189 bp band on gels after electrophoresis of the PCR products. BAL: bronchoalveolar lavage; ICAM-1: intercellular adhesion molecule-1; mFI: mean fluorescence; mRNA: messenger ribonucleic acid; ND: not determined.

ture in 5 of the 6 samples, but at a low level. The level of ICAM-1 mRNA was evaluated according to the optical density (OD) of the 189 bp band for each sample. AM culture with rIFN- γ induced ICAM-1 mRNA expression in every case ($p < 0.01$, OD before culture *vs* OD after 24 h of culture) and OD values correlated with mFI values ($r = 0.73$; $n = 6$) but not with the percentage of CD54+ cells ($r = 0.37$). ICAM-1 mRNA synthesis was significantly inhibited by 8 $\mu\text{g}\cdot\text{mL}^{-1}$ fusafungine ($p < 0.05$, OD at 8 $\mu\text{g}\cdot\text{mL}^{-1}$ *vs* OD at 0 $\mu\text{g}\cdot\text{mL}^{-1}$). At 2 $\mu\text{g}\cdot\text{mL}^{-1}$ fusafungine, ICAM-1 mRNA expression was inhibited in 3 of the 6 samples. Figure 3 illustrates typical results of ICAM-1 antigen analysis by means of immunocytofluorimetry and that of ICAM-1 mRNA using PCR, before culture and after 24 h of culture with rIFN- γ and various concentrations of fusafungine. In this experiment, ICAM-1 mRNA transcription (expressed as OD) fell as the concentration of fusafungine increased. Surprisingly, GAPDH mRNA levels fell in parallel. However, whilst the ICAM-1 mRNA OD was four times lower with 16 $\mu\text{g}\cdot\text{mL}^{-1}$ fusafungine than without fusafungine, the GAPDH mRNA OD only fell by 1.8 times.

Fusafungine, thus, significantly inhibited ICAM-1 mRNA expression. However, the inhibitory effect on ICAM-1 transcription was observed at higher concentrations than those inhibiting ICAM-1 antigen expression, suggesting that fusafungine acts mainly on post-transcription rather than transcription.

Persistent effect of fusafungine on ICAM-1 expression

Whether the inhibitory effect of fusafungine persisted after drug removal was then investigated, on one

BAL containing a sufficient number of AMs. After 24 h of culture with rIFN- γ and fusafungine, supernatants were removed and fresh supplemented RPMI medium containing no rIFN- γ or fusafungine was added. After a further 24 h of incubation, AMs were analysed for ICAM-1 antigen and mRNA expression. Fusafungine inhibited ICAM-1 expression at both the post-transcriptional and transcriptional levels, in a concentration-dependent manner (fig. 4). After removal both of rIFN- γ and fusafungine, the inhibitory effect previously observed at 16 $\mu\text{g}\cdot\text{mL}^{-1}$ fusafungine on the percentage of ICAM-1-positive cells and on mRNA synthesis persisted. However, antigen density was higher, probably owing to the time lag between mRNA transcription and translation. At a lower concentration of fusafungine (2 $\mu\text{g}\cdot\text{mL}^{-1}$), the stimulatory effect of rIFN- γ on ICAM-1 expression persisted, but not the inhibitory effect of fusafungine.

These experiments indicated that fusafungine inhibited the rIFN- γ -induced expression both of ICAM-1 protein and mRNA. Moreover, the inhibitory effect observed with high doses of fusafungine persisted after removal both of the drug and the stimulator.

Comparative effects of fusafungine on spontaneous and rIFN- γ -induced ICAM-1 expression by AMs

The effects of fusafungine on basal and rIFN- γ -induced ICAM-1 expression by freshly isolated human AMs were then compared. Before culture, 20% of BAL cells were CD54-positive, with an mFI of 65 (fig. 5). Culture in the presence or absence of rIFN- γ resulted in increased ICAM-1 antigen expression (88 and 86% of CD54-positive cells, respectively) but at a low level in the absence








time hr		0		24			
Fusafungine $\mu\text{g}\cdot\text{mL}^{-1}$		0	0	2	8	16	
% CD54 positive cells		34	94	94	28	39	
mFI		43	231	235	140	146	
ICAM-1	189 bp OD						
		0	0.29	0.48	0.37	0.12	
GAPDH	574 bp OD						
		1.11	1.42	1.52	1.29	0.81	

Fig. 3. - Effect of fusafungine on ICAM-1 expression by alveolar macrophages from one representative BAL sample after 24 hr culture with 250 U·mL⁻¹ rIFN- γ . ICAM-1 protein expression was determined by immunocytofluorimetric analysis using the CD54 MoAb. Results were recorded as the percentage of ICAM-1 positive cells and the mean fluorescence intensity. ICAM-1 mRNA was determined by PCR on cDNA obtained after specific reverse transcription. The PCR products were evaluated by densitometric analysis of the expected 189 bp-band on gel electrophoresis and the results expressed in arbitrary units of optical density (OD). GAPDH mRNA was determined in parallel as control. bp: base pairs; OD: optical density; GAPDH: glyceraldehyde-3-phosphate dehydrogenase. For further abbreviations see legend to figure 1.

time hr	IFN + fusafungine						New medium without IFN or fusafungine		
	0	0	2	16			0	2	16
Fusafungine $\mu\text{g}\cdot\text{mL}^{-1}$	0	0	2	16			0	2	16
% CD54 positive cells	11	83	50	31			84	83	30
mFl	52	80	69	92			195	150	162
ICAM-1 189 bp OD									

Fig. 4. – Persistence of the inhibitory effect of fusafungine on rIFN- γ -induced ICAM-1 expression by alveolar macrophages from one representative BAL. AMs were cultured for 24 hr with rIFN- γ in the presence or absence of fusafungine; the medium was then replaced and cells were further incubated for 24 hr without rIFN- γ or fusafungine. Protein expression was determined by immunocytofluorimetric analysis using the CD54 MoAb. Results were recorded as the percentage of ICAM-1 positive cells and the mean fluorescence intensity. ICAM-1 mRNA was determined by PCR on cDNA obtained after specific reverse transcription. The PCR products were evaluated by densitometric analysis of the expected 189 bp-band on gel electrophoresis and the results expressed in arbitrary units of optical density (OD). For abbreviations see legends to figure 1 and 3.

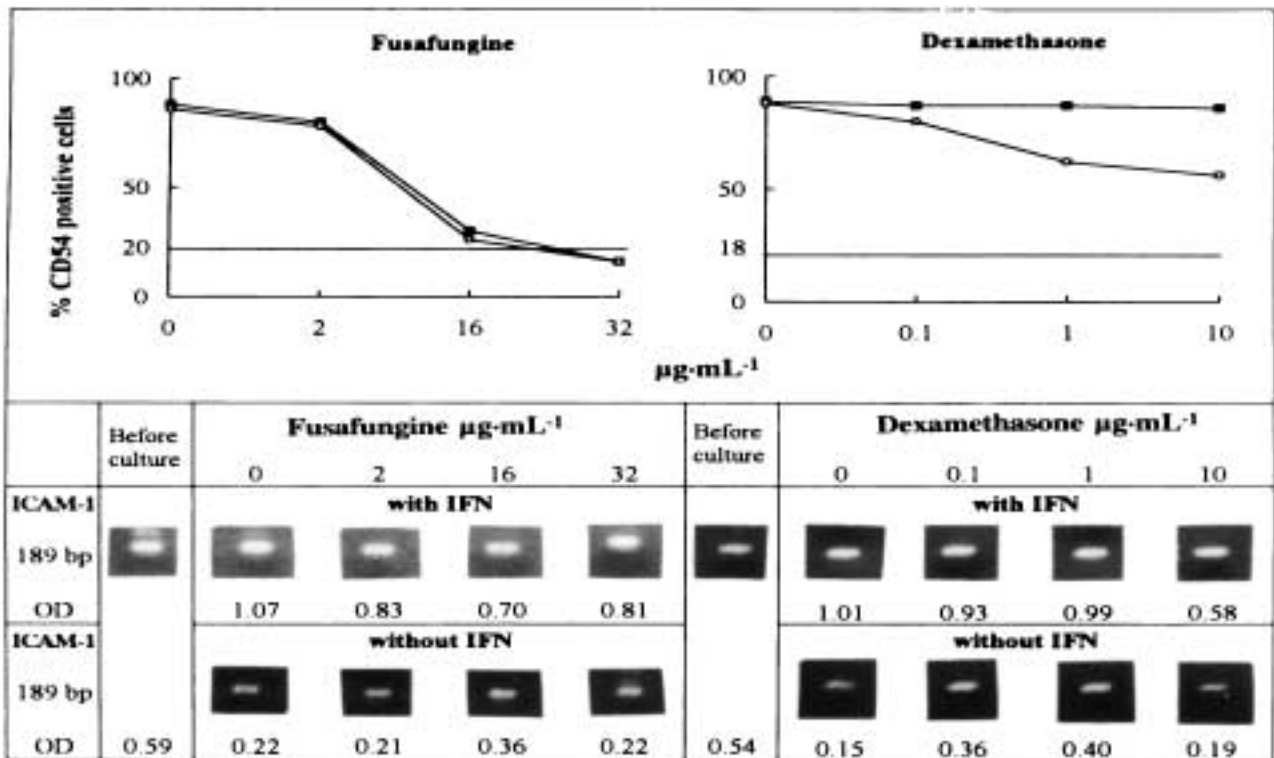


Fig. 5. – Comparative effects of fusafungine and dexamethasone on spontaneous and rIFN- γ -induced ICAM-1 expression by alveolar macrophages after 24 h culture. Upper panels: ICAM-1 protein expression was determined as the percentage of ICAM-1-positive cells in flow cytometry with anti-CD54 MoAb. Culture in the presence (—■—) or absence (—○—) of 250 U·mL⁻¹ rIFN- γ . The horizontal line represents the percentage of CD54-positive cells before culture. Lower panel: ICAM-1 mRNA was determined by PCR on cDNA obtained after specific reverse transcription. The PCR products were evaluated by densitometric analysis of the expected 189 bp band on gel electrophoresis and the results are expressed in arbitrary optical density (OD) units. mRNA: messenger ribonucleic acid; PCR: polymerase chain reaction; cDNA: complementary deoxyribonucleic acid. For further abbreviations see legends to table 1 and 3.

of rIFN- γ induction (mFI=277 and 108, respectively). Surprisingly, ICAM-1 mRNA expression increased only after rIFN- γ induction (OD 1.07 after culture and 0.59 before culture), suggesting that the enhanced ICAM-1 protein expression observed after culture in the absence of rIFN- γ only resulted from post-transcriptional upregulation, whereas an additional transcriptional upregulation was observed in the presence of rIFN- γ .

As shown in figure 5, fusafungine inhibited both spontaneous and rIFN- γ -induced ICAM-1 protein expression on BAL cells in a concentration-dependent manner. ICAM-1 mRNA expression was slightly inhibited in rIFN- γ -treated cells only, whatever the concentration of fusafungine, indicating that fusafungine antagonizes the stimulatory effect of rIFN- γ on ICAM-1 protein expression and downregulates spontaneous expression in a concentration-dependent manner.

Dexamethasone inhibits basal but not IFN- γ -induced ICAM-1 expression on AM in vitro

Finally, the effects of fusafungine, a nonsteroidal anti-inflammatory drug, on ICAM-1 expression in human BAL cells were compared with that of a glucocorticoid, dexamethasone. BAL cells were treated for 24 h with 0, 0.1, 1 and 10 $\mu\text{g}\cdot\text{mL}^{-1}$ of dexamethasone (2.5×10^{-4} to 2.5×10^{-2} M), in the presence and absence of rIFN- γ . Before culture, 18% of AMs were CD54-positive with an mFI of 47, confirming the basal membrane expression of ICAM-1. ICAM-1 expression was markedly increased in the presence of rIFN- γ , as the percentage of CD54-positive cells reached 89% after 24 h of culture and as antigen density was high (mFI=320). ICAM-1 expression was also enhanced in the absence of rIFN- γ , as the percentage of CD54-positive cells reached 88%; however, antigen density was lower (mFI=116). As observed in the experiments with fusafungine, ICAM-1 mRNA levels increased only after rIFN- γ induction (OD after culture 1.01, OD before culture 0.54), whereas it fell after culture without rIFN- γ (OD=0.15). This suggested that the enhanced ICAM-1-protein expression observed after culture without rIFN- γ resulted only from post-transcriptional upregulation, whereas culture with rIFN- γ led to an additional transcriptional upregulation.

Dexamethasone did not influence rIFN- γ -induced ICAM-1 expression on AMs (fig. 5), even at the highest concentration (10 $\mu\text{g}\cdot\text{mL}^{-1}$). Only the intensity of ICAM-1 expression on the AM surface was slightly reduced. The OD of the 189 bp ICAM-1 band following PCR on cDNA was reduced at 10 $\mu\text{g}\cdot\text{mL}^{-1}$ only and correlated with neither the percentage of CD54-positive cells nor the mFI. In the absence of rIFN- γ , dexamethasone partially inhibited ICAM-1 expression on AMs in a concentration-dependent manner, as the percentage of positive cells fell to 56% at 10 $\mu\text{g}\cdot\text{mL}^{-1}$, whereas the mFI remained steady whatever the dexamethasone concentration. Surprisingly, ICAM-1 mRNA levels were enhanced by culture without rIFN- γ in the presence of 0.1–1 $\mu\text{g}\cdot\text{mL}^{-1}$ dexamethasone, but not by 10 $\mu\text{g}\cdot\text{mL}^{-1}$. These results indicate that high concentrations of dexa-

methasone antagonize the stimulatory effect of rIFN- γ on ICAM-1 mRNA expression on BAL cells without reducing ICAM-1 protein levels. In contrast, dexamethasone downregulated basal expression of ICAM-1 antigen on BAL cells without affecting mRNA levels. Note that dexamethasone had no effect on HLA-DR expression.

Discussion

ICAM-1 expression on alveolar macrophages is increased in inflammatory lung diseases, and *in vitro* studies have indicated that ICAM-1 expression is regulated by rIFN- γ and IL-1 produced by activated T-lymphocytes and macrophages [20]. It seems reasonable to assume that downregulation of ICAM-1 expression on AMs in airway tissues by drugs may constitute a therapeutic mechanism in the lung inflammatory process. Most previous studies of ICAM-1 regulation on monocytes/macrophages have involved rat thymic macrophages [9], human blood monocytes [11], monocytic cell lines [10], or macrophage cell lines [21]. In the present study, we compared the effects of fusafungine and dexamethasone on ICAM-1 expression in human AMs freshly recovered from 14 BAL samples.

Treatment of AMs with rIFN- γ for 24 h induced a significant increase in ICAM-1 expression, mainly at the transcriptional level. Culture without rIFN- γ also resulted in a marked increase in the percentage of AMs expressing ICAM-1 protein, but at a lower intensity. Here ICAM-1 upregulation appeared to occur at a post-transcriptional level and might have involved activated migration of the intracytoplasmic protein to the cell membrane. Our results indicate that rIFN- γ is a potent inducer of ICAM-1 expression in AMs, in contrast to lipopolysaccharide (LPS) (data not shown).

As regards the mechanism of ICAM-1 downregulation *in vitro*, fusafungine may act at the transcriptional or post-transcriptional level, or inhibit protein migration to the cell surface. Fusafungine ($>8 \mu\text{g}\cdot\text{mL}^{-1}$) inhibited the stimulatory effect of rIFN- γ on ICAM-1 mRNA expression in a concentration-dependent manner. However, GAPDH mRNA levels also fell as the fusafungine concentration increased, meaning that we could not quantify ICAM-1 mRNA relative to GAPDH mRNA. Fusafungine also downregulated β -actin mRNA (data not shown). Since the downregulation of GAPDH mRNA by fusafungine was 2.2 times weaker than that of ICAM-1 mRNA at a given concentration, fusafungine acts more specifically on ICAM-1 transcription gene than on others. Fusafungine ($>2 \mu\text{g}\cdot\text{mL}^{-1}$) inhibited rIFN- γ -induced ICAM-1-antigen expression in a concentration-dependent manner, reaching a total inhibition at 16 $\mu\text{g}\cdot\text{mL}^{-1}$. ICAM-1 antigen density fell in parallel. However, the expression of other membrane antigens, such as HLA-DR, was not altered. Altogether, these data strongly suggest that the marked inhibitory effect of low concentrations of fusafungine on rIFN- γ -induced ICAM-1 expression by AMs acts at a post-transcriptional level, whereas high concentrations of the drug act on the transcription of numerous genes, although more markedly on ICAM-1 gene.

The mechanism by which fusafungine regulates ICAM-1 expression may be related, at least in part, to its ionophoric properties, as the drug acts as a mobile cation carrier at concentrations higher than $0.01 \mu\text{g}\cdot\text{mL}^{-1}$. Fusafungine is a mixture of seven enniatins, *i.e.* cyclohexadepsipeptides with ionophoric properties, that modify membrane permeability to potassium. Because of their high solubility in polar media, enniatins enter the inner cell compartment, thus directly influencing basic vital processes [22, 23]. Moreover, their ability to fix and transport through cell membranes not only K^+ but also Na^+ and divalent cations, such as Mg^{2+} and Ca^{2+} , might be responsible for alterations of the vital functions, such as the overall transcription process.

Why the inhibitory effect of fusafungine at high concentrations persisted after drug removal may be explained by the fact that inhibition of ICAM-1 mRNA synthesis by high concentrations of the drug is only overcome after at least 24 h of culture in the absence of the drug. Alternatively, fusafungine may persist inside the cells after drug removal. In contrast, at low concentrations fusafungine did not inhibit rIFN- γ -mediated activation that persisted following removal of the cytokine.

ICAM-1 appears to play an important role in the inflammatory response during lung allograft rejection. ICAM-1 downregulation may explain the effects of treatments for acute rejection. Indeed, inhibition of lymphocytic and neutrophilic cell infiltration of perivascular spaces increases with the intensity of immunosuppressive treatment in transbronchial lung biopsies during acute lung rejection [24]. Dexamethasone rapidly acts on pulmonary inflammation characterized by APC and T-cell activation, T-cell proliferation and cytotoxic functions, suggesting that it might act on cellular adhesion mechanisms. Our findings provide evidence that the anti-inflammatory glucocorticoid dexamethasone, unlike fusafungine, may not antagonize rIFN- γ -induced ICAM-1 protein expression, despite a partial inhibitory effect on ICAM-1 gene transcription at a high concentration. In culture, dexamethasone appeared to partially inhibit spontaneous ICAM-1-protein expression but not ICAM-1-gene transcription.

These results are in keeping with reports that 10^{-5} M dexamethasone does not abrogate TNF-induced expression of ICAM-1 on human dermal microvascular endothelial cells (HDMEC), as shown by means of immunocytochemistry [25]. More recently, the same authors tested four steroids for their effects on HDMEC treated for 24 h with TNF and/or IL-1 β , which appear to be strong inducers of ICAM-1 expression. None of the steroids, which included dexamethasone, influenced cytokine-induced ICAM-1 expression, strongly suggesting that the anti-inflammatory effects of steroids are not related to ICAM-1 modulation, at least in HDMEC [26]. In contrast, using immunoperoxidase-based techniques, other authors found that ICAM-1 expression was upregulated on freshly isolated rat thymic macrophages after culture with rIFN- γ , IL-1 and TNF- γ , and downregulated by dexamethasone [9]. Again, dexamethasone inhibits both constitutive and IL-3- or GM-CSF-induced expression of the ICAM-1 antigen on purified normal human blood

monocytes [11]. It has also been reported that dexamethasone can inhibit ICAM-1 transcription directly both in undifferentiated and differentiated (macrophage-like) cells of the human monocytic line U937, as shown by means of flow cytometry and northern blotting [10]. In our model, dexamethasone appeared to downregulate spontaneous ICAM-1 protein expression on human AM in culture and to antagonize the stimulatory effect of rIFN- γ on ICAM-1 transcription. Fusafungine antagonized the stimulatory effect of rIFN- γ on ICAM-1 protein expression and downregulated its spontaneous expression, but did not appear to act directly at the transcriptional level.

Drugs like fusafungine that locally inhibit ICAM-1 expression by AMs may have therapeutic potential in inflammatory diseases of the respiratory tract. The observation that dexamethasone did not inhibit rIFN- γ -induced ICAM-1 expression in our human AM model strongly suggests that anti-inflammatory drugs act on different cellular and molecular targets and should, thus, be studied in various *in vitro* systems.

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