Characterization of the pharmacological profile of the potent LTB₄ antagonist CP-105,696 on murine LTB₄ receptors *in vitro*

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1 Binding of [³H]-leukotriene B₄ ([³H]-LTB₄) to murine spleen membranes (MSM) was determined.

2 Scatchard analyses of $[^{3}H]$ -LTB₄ binding indicated the presence of high $(K_{D1} = 1.7 \text{ nM})$ and low $(K_{D2} = 7.5 \text{ nM})$ affinity receptors on MSM with B_{max} values of 151 fmol mg⁻¹ protein (B_{max1}) and 354 fmol mg⁻¹ protein (B_{max2}) , respectively.

3 CP-105,696, a potent LTB₄ antagonist, inhibited [³H]-LTB₄ (0.67 nM) binding to the high affinity receptor on MSM, IC_{50} = 30.2 nM, K_i = 17.7 nM with a Hill coefficient of 0.93.

4 Scatchard analyses of $[^{3}H]$ -LTB₄ binding to MSM in the presence of CP-105,696 indicated that the high-affinity receptor was inhibited in a non-competitive manner and the low-affinity receptor in a competitive manner.

5 Isolated peripheral blood murine neutrophils (MN) responded chemotactically to LTB_4 , $EC_{50} = 2.5$ nM. CP-105,696 blocked this response, $IC_{50} = 2.3$ nM. When examined over a full concentration-response range of LTB_4 , CP-105,696 inhibited chemotaxis in a non-competitive manner. 6 Murine neutrophils in anticoagulated whole blood upregulated the integrin, complement receptor type 3 (CD11b/CD18, Mac-1) in response to LTB_4 , $EC_{50} = 20$ nM and this was inhibited by CP-105,696 in a competitive manner.

7 These results provide evidence that MSM have specific binding sites for LTB₄, and as exemplified by CP-105,696, that these receptors may be useful for determining the potency and nature of antagonism of novel LTB₄ receptor antagonists.

Keywords: Leukotriene B₄ receptors; murine spleen membranes; neutrophils; chemotaxis; complement receptor 3; CD11b/ CD18; Mac-1; CP-105,696

Introduction

Leukotriene B₄, a product of the 5-lipoxygenase pathway of arachidonic acid metabolism (Borgeat & Samuelsson, 1979) is a potent proinflammatory mediator thought to be involved in the pathogenesis of several inflammatory diseases (Ford-Hutchinson, 1990). Many reports over the past decade have provided evidence of specific high affinity receptors for LTB₄ on inflammatory cells such as neutrophils eosinophils and macrophages from several different species (Goldman & Goetzl, 1984; Lin et al., 1984; Cristol et al., 1988; Maghni et al., 1991). In addition, membranes derived from guinea-pig and porcine spleen have also been shown to bind LTB_4 with high affinity (Cheng et al., 1986; Miki et al., 1990). Although there are a number of published reports demonstrating LTB₄mediated leukocyte infiltration in vivo in mice (Suarez et al., 1987; Thieroff et al., 1988; Fretland et al., 1990; Griswold et al., 1991; Iwamoto et al., 1993) to our knowledge there are no published reports showing direct evidence of LTB₄ receptors on murine cells or tissue. We therefore undertook the series of experiments described herein to determine if LTB₄ receptors could be detected on murine tissue (e.g. spleen) and to see if evidence could be provided that would link these receptors to functional responses of murine peripheral blood neutrophils. The ultimate goal of these experiments was to determine if the novel LTB₄ antagonist, CP-105,696 (Koch et al., 1994) maintained potency on murine tissue/cells that would support its in vivo evaluation in murine inflammatory disease models.

Methods

Membrane preparation

Spleens were dissected from 40 to 50 BALB/c mice and kept in 50 mM Tris-HC1 buffer (pH=7.3 at 25°C). After removal of connective tissue, the total weight of the spleens was determined. The pooled tissue was minced and homogenized twice by Polytron (Brinkmann Instruments, Westbury, NY, U.S.A.) PCU-2 at setting 7 for 10 s in 50 mM Tris at 3 ml g⁻¹ tissue. The homogenate was centrifuged at 1000 g for 15 min and the resultant supernatant was centrifuged at 40,000 g for 20 min with washing of the intermediate pellet with 50 mM Tris. The final pellet was resuspended at a protein concentration of 5 mg ml⁻¹ in 50 mM Tris pH=7.3 with 10 mM MgCl₂ and stored at -70° C. The protein concentration was determined by the Bradford method (Bradford, 1976) using bovine serum albumin as standard.

$[^{3}H]$ -LTB₄ receptor binding assays

The [³H]-LTB₄ mouse spleen binding assay was performed according to a method adapted from Cheng *et al.* (1986). The [³H]-LTB₄ binding assays were carried out in a volume of 150 μ l containing 50 μ g membrane preparation, 0.67 nM [³H]-LTB₄, 50 mM Tris pH = 7.3, 10 mM MgCl₂ and 10% methanol in the presence or absence of various concentrations of unlabelled LTB₄ or CP-105,696 for 30 min at 4°C. The assay was performed in triplicate in microtitre plates (Costar, Cambridge, MA, U.S.A.) and filtered through Whatman GF/B glass fibre filters using a beta plate apparatus (Pharmacia LKB, Piscataway, NJ, U.S.A.) to separate free from bound [³H]-LTB₄. Specific [³H]-LTB₄ binding represents the value obtained when nonspecific binding (value obtained in the

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presence of 5 μ M unlabelled LTB₄) is subtracted from total binding. For saturation binding experiments, concentrations of [³H]-LTB₄ ranging from 0.25 to 20 nM were used. [³H]-LTB₄ binding to the spleen membranes was unchanged for up to 6 months.

Murine neutrophil isolation and chemotaxis assay

Neutrophils were isolated from pooled anticoagulated blood obtained from BALB/c mice by intracardiac puncture according to the procedure described by Rot (1991). Isolated cells (usually $\sim 30\%$ neutrophils) were adjusted to a density of 2.5×10^6 neutrophils ml⁻¹ in HBSS containing 1.6 mM Ca²⁺ and 0.7 mM Mg²⁺, 10 mM HEPES and 1% BSA (pH 7.2). Chemotaxis was performed using a 48-well chamber apparatus (Neuroprobe, Cabin John, MD, U.S.A.) with cellulose nitrate filters (pore size, 3.0 μ m) as previously described (Harvath et al., 1987). Under these assay conditions, neutrophils are the only cells that migrate into the filters. The total number of cells (observed at $\times 400$ magnification in 3-5 fields) migrating from 20 μ m beneath the surface monolayer to the leading front (usually $40-60 \ \mu m \ 60 \ min^{-1}$ at optimal chemotactic factor concentrations) were summed and provided an index of the chemotactic response. Each experiment was performed on at least three separate occasions. In experiments where CP-105,696 was examined for antagonist properties, it was present in both compartments of the chemotaxis chamber.

Upregulation of Mac-1 on neutrophils in whole blood

Whole blood obtained by intracardiac puncture from BALB/c mice was anticoagulated with EDTA (0.37%); 100 μ l of blood was then added to 12×75 mm borosilicate test tubes containing 10 μ l of 10 × stock concentration of CP-105,696 for 5 min at 37°C. LTB₄ (20 nM final concentration) was added to the samples and incubated for 10 min at 37°C. Tubes were then removed and placed in an ice water bath and ~ 1 ml of cold PBS containing 10 mM EDTA, pH 7.25, 0.2% sodium azide and 0.2% heat-inactivated foetal bovine serum (PBS-wash). The cells were then pelleted at 200 g and incubated with either FITC-conjugated rat anti-mouse Mac-1 mAb or rat IgG-2b isotype control Ab for the mouse cells. The RBC were lysed with FACS Lysing Solution (Becton-Dickinson, San Jose, CA, U.S.A.) and the remaining cells washed with PBS-wash at 4°C. Fluorescent labelling was quantified immediately with a B.D. FACScan flow cytometer with a live gate on the neutrophils. Data from 5000 neutrophils per sample were recorded and expressed as the percentage maximal mean channel fluorescence.

Data analysis

Results are expressed as the means \pm s.e.mean from at least three independent experiments unless otherwise indicated. IC₅₀ values were calculated by plotting the mean of the percent control of each compound-treated group on a logscale, fitting the linear portion of the graphs to a linear regression curve and calculating the x value at y=50. Binding site densities and respective affinities for [³H]-LTB₄ were calculated according to the published method (Scatchard, 1949) using London Software Receptor Fit Saturation Two-site Program. K_i was calculated according to the method of Cheng & Prusoff, 1973.

Materials

[³H]-LTB₄ (specific activity = 200 Ci mmol⁻¹) was purchased from DuPont/NEN (Boston, MA, U.S.A.); LTB₄ from Biomol, Plymouth Meeting, PA, U.S.A., BALB/c mice from Jackson Laboratories, Bar Harbor, ME, U.S.A. CP-105,696, (+)-1-(3**S**,4**R**)-[3-(4-phenyl-benzyl)-4-hydroxy-chroman-7-yl]cyclopentane carboxylic acid was synthesized according to Koch *et al.* (1994) and its structure is shown in Figure 1.

Results

Evidence for specific $[^{3}H]$ -LTB₄ binding sites on MSM

Figure 2 shows a representative saturation isotherm and Scatchard plot for the binding of $[{}^{3}H]$ -LTB₄ to murine spleen membranes. Analysis of this data indicated the presence of two distinct binding sites with the higher affinity site exhibiting a dissociation constant (K_{D1}) (mean ± s.e.) of 1.7 ± 0.4 nM (n=3) and a total binding capacity (B_{max1}) of 151 ± 17 fmol mg⁻¹ protein, and a lower affinity binding site exhibiting a dissociation constant (K_{D2}) of 7.5 ± 2.1 nM (n=3) and a total binding capacity (B_{max2}) of 354 ± 59 fmol mg⁻¹ protein. [${}^{3}H$]-LTB₄ binding was linear over a protein concentration range of $5-50 \mu g$ /well and half-maximal binding occurred at ~ 10 min reaching equilibrium at 30 min (results not shown).

Effect of CP-105,696 on $[^{3}H]$ -LTB₄ binding to high affinity receptors on MSM

Figure 3 demonstrates the ability of CP-105,696 to compete for $[{}^{3}H]$ -LTB₄ binding to the high-affinity binding site on MSM. An IC₅₀ value of 30.2 ± 1.2 nM ($K_{i} = 17.7 \pm 0.7$ nM) (n = 3) was obtained for CP-105,696; this is ~10 fold higher than the IC₅₀ value obtained for LTB₄, 3.1 ± 0.5 nM (n = 3). Scatchard analysis of $[{}^{3}H]$ -LTB₄ binding to high-affinity receptors on MSM in the presence and absence of fixed concentrations of CP-105,696 showed clear evidence of noncompetitive interactions of CP-105,696 on $[{}^{3}H]$ -LTB₄ binding to this high affinity site (Figure 4). The $K_{\rm D}$ values of $[{}^{3}H]$ -LTB₄ obtained in the presence of CP-105,696 at 1 nM and 5 nM were 1.9 nM and 1.9 nM



Figure 1 Structure of CP-105,696.



Figure 2 Binding of increasing concentrations of $[{}^{3}\text{H}]$ -LTB₄ to MSM (inset) and Scatchard plot of receptor binding data. Data shown are representative of three independent experiments. Data were fitted to a two-site model using the Lundon Software Receptor Fit Saturation Binding Programme. Symbols in inset represent (\bigcirc) total, (\blacksquare) nonspecific and (\bigcirc) specific $[{}^{3}\text{H}]$ -LTB₄ binding. Calculated $K_{\text{D1}} = 1.98$ nM and $B_{\text{max1}} = 118$ fmolmg⁻¹ (\bigcirc) and $K_{\text{D2}} = 8.98$ nM and $B_{\text{max2}} = 270$ fmol mg⁻¹ (\bigcirc).

respectively compared to the control value of 2.0 nM. The corresponding densities of high-affinity [³H]-LTB₄ binding sites per cell in the presence of CP-105,696 at 1.0 nM and 5.0 nM were 105 fmol mg⁻¹ and 76 fmol mg⁻¹ respectively, compared with the control value of 118 fmol mg⁻¹.

Effect of CP-105,696 on $[^{3}H]$ -LTB₄ binding to lowaffinity receptors on MSM

Figure 5 shows the Scatchard analysis of $[{}^{3}H]$ -LTB₄ binding to low-affinity receptors on MSM in the presence and absence of fixed concentrations of CP-105,696. Evidence of competitive interactions at this low-affinity binding site were obtained. The $K_{\rm D}$ values of $[{}^{3}H]$ -LTB₄ obtained in the presence of CP-105,696 at 1 nM and 5 nM were 16 nM and 22 nM respectively compared to the control value of 10.0 nM. The corresponding

100 80 maximum specific [³H-LTB₄ binding 60 40 % 20 لسبب 0 10⁴ 0.1 1 10 100 1000 Competing ligand concentration (пм)

Figure 3 Inhibition of $[{}^{3}H]$ -LTB₄ binding to MSM by CP-105,696 (O) and LTB₄ ($\textcircledoldsymbol{\bullet}$). IC₅₀ for LTB₄=3.14±0.48 nM and for CP-105,696=30.2±1.2 nM. MSM were incubated with 0.67 nM $[{}^{3}H]$ -LTB₄ for 30 min at 4°C in the presence and absence of various concentrations of CP-105,696 or LTB₄ and specific binding was determined. Data are means±s.e. from three experiments each performed in triplicate. Zero percent specific binding refers to the level of $[{}^{3}H]$ -LTB₄ binding obtained in the presence of 1.0 μ M unlabelled LTB₄.



Figure 4 Scatchard plots of $[^{3}H]$ -LTB₄ binding to high-affinity receptors on MSM in the absence (\bigoplus) and presence of (\bigcirc) 1 nM and (\bigoplus) 5 nM CP-105,696. MSM were incubated with concentrations of $[^{3}H]$ -LTB₄ (0.25 nM-2.0 nM) for 30 min at 4°C in the absence and presence of CP-105,696. Control $K_{D} = 1.98$ nM, $B_{max} = 118$ fmol mg⁻¹; 1 nM CP-105,596, $K_{D} = 1.93$ nM, $B_{max} = 105$ fmol mg⁻¹; 5 nM CP-105,596, $K_{D} = 1.93$ nM, $B_{max} = 105$ fmol mg⁻¹; 5 nM CP-105,596, $K_{D} = 1.85$ nM, $B_{max} = 76.3$ fmol mg⁻¹. Data shown are representative of three independent experiments. Data were fitted to a one-site model using the Lundon Software Receptor Fit Saturation Binding Programme.

densities of low affinity [³H]-LTB₄ binding sites per cell in the presence of CP-105,696 at 1.0 nM and 5.0 nM were 456 fmol mg⁻¹ and 466 fmol mg⁻¹ respectively, compared with the control value of 468 fmol mg⁻¹.

Effect of CP-105,696 on LTB₄-mediated murine neutrophil chemotaxis

Using the 48-microwell chemotaxis chamber and cellulose nitrate filters of 3 μ m pore size we observed half-maximal effects of LTB₄ on neutrophil migration at 2.5 nM and maximal effects at 100 nM (see inset Figure 6). With 5 nM as the fixed concentration of LTB₄ in the lower chamber, half maximal inhibition of chemotaxis occurred at a concentration of 2.3±0.7 nM (n=5) of CP-105,696. When chemotaxis was assessed over a full range of concentrations of LTB₄, the effect of



Figure 5 Scatchard plots of $[{}^{3}H]$ -LTB₄ binding to low-affinity receptors in MSM in the absence (•) and presence of (○) 1 nM and (•) 5 nM CP-105,696. MSM were incubated with concentrations of $[{}^{3}H]$ -LTB₄ (4.0 nM - 20.0 nM) for 30 min at 4°C in the absence and presence of CP-105,696. Control, $K_{D} = 10.2$ nM, $B_{max} = 468$ fmol mg⁻¹; 1 nM CP-105,596, $K_{D} = 16.0$ nM, $B_{max} = 456$ fmol mg⁻¹; 5 nM CP-105,596, $K_{D} = 21.8$ nM, $B_{max} = 466$ fmol mg⁻¹. Data shown are representative of three independent experiments. Data were fitted to a one-site model using the Lundon Software Receptor Fit Saturation Binding Programme.



Figure 6 Inhibition of LTB₄-mediated murine neutrophil chemotaxis by CP-105,696. LTB₄ was used at a fixed concentration of 5 nM in the lower chamber and CP-105,696 was present at various concentrations $(10 \text{ pM} - 1 \mu\text{M})$ in both the upper and the lower chamber. Chambers were incubated for 60 min at 37°C and then the filters were removed, processed and counted as described in Methods. Data are the means \pm s.e. from 5 independent experiments. Inset shows LTB₄ concentration-response curve.



Figure 7 Effect of CP-105,696 on the concentration-dependent curve of LTB₄-mediated chemotaxis. Chemotaxis was induced by a range of concentrations of LTB₄ ($10 \text{ pm} - 1 \mu \text{M}$) in the absence (\bigcirc) and presence of (\triangle) 10^{-9} M; (\triangle) 10^{-8} M; (\square) 10^{-7} M; (\blacksquare) 10^{-6} M; (\bigcirc) 10^{-5} M CP-105,696. Data are expressed as the percentage of maximal chemotaxis occurring to LTB₄ in the absence of CP-105,696 and are means ± s.e. from three independent experiments.



Figure 8 Inhibition of LTB₄-mediated Mac-1 upregulation in murine neutrophils in whole blood. LTB₄ was used at a fixed concentration of 100 nm. CP-105,696 was present at various concentrations $(35-140 \,\mu\text{M})$. Data are expressed as the percentage of the response occurring in the absence of CP-105,696 and are the means \pm s.e. from 4 independent experiments. Inset shows LTB₄

the addition of CP-105,696 could not be overcome by increasing concentrations of LTB₄, which offers evidence that a noncompetitive form of inhibition is occurring at the LTB₄ receptor mediating neutrophil chemotaxis (Figure 7).

Effect of CP-105,696 on LTB_4 -mediated upregulation of the adhesion molecule Mac-1 on murine neutrophils in whole blood

Half-maximal effects of LTB₄ on Mac-1 upregulation on murine neutrophils in whole blood occurred at ~20 nM with maximal effects at ~1.0 μ M (see inset Figure 8). Using 100 nM as the fixed concentration of LTB₄, we observed half maximal inhibition of Mac-1 upregulation by CP-105,696 at 66.5±10.7 μ M (n=4). When Mac-1 upregulation was examined over a full range of concentrations of LTB₄, the addi-



Figure 9 Effect of CP-105,696 on the concentration-dependent curves of LTB₄-mediated upregulation of Mac-1 in neutrophils in whole blood. Mac-1 upregulation was induced by a range of concentrations of LTB₄ in the absence (\bullet) and presence of (\bigcirc) 35 μ M; (\square) 70 μ M; (\square) 140 μ M CP-105,696. Data are expressed as the percentage of maximal response occurring to LTB₄ in the absence of CP-105,696 and are from four experiments.

tion of increasing concentrations of CP-105,696 caused parallel rightward shifts in the LTB_4 -concentration curves, which indicates that competitive interactions at the LTB_4 receptor mediating CD11b upregulation are occurring (Figure 9).

Discussion

Several published reports have provided evidence of specific LTB₄ receptors on inflammatory cells and that these receptors were probably linked to functional activation of cells such as human neutrophils (Goldman & Goetzl, 1984; Lin et al., 1984). More recently through the use of the specific LTB₄ receptor antagonist, CP-105,696, we have presented pharmacological evidence that the high affinity LTB₄ receptor ($K_D = 0.65$ nM) appears to mediate chemotaxis, and the low affinity LTB₄ receptor ($K_D = 5.8$ nM) is linked to secretory events in human neutrophils (Showell et al., 1995). Two other reports have demonstrated the presence of high affinity binding sites for LTB₄ on membranes derived from guinea-pig and porcine spleen (Cheng et al., 1986; Miki et al., 1990). Although the specific cell source(s) contributing these binding sites for [³H]-LTB₄ were not defined it was felt that these receptors were physiologically relevant, and for guinea-pigs, possibly unrelated to the LTB₄ binding site present on granulocytes from this species (Cheng et al., 1986). As it is impractical to obtain sufficient numbers of circulating murine neutrophils suitable for [³H]-LTB₄ receptor binding assays and as mouse spleen is known to serve a myelopoietic function (Bannerman, 1983) we reasoned that membranes derived from this tissue may be a useful surrogate for murine neutrophils and thus possess LTB₄ receptors reflective of this cell population. Using [³H]-LTB₄ as a specific probe we show here evidence for the presence of specific binding sites for LTB4 on membranes derived from murine spleen with affinities in the nanomolar range. Scatchard analysis of the [3H]-LTB4 binding isotherms revealed the presence of two distinct receptors with affinities of 1.7 nM (K_{D1}) and 7.5 nm (K_{D2}) respectively. Interestingly, guinea-pig (Cheng et al., 1986) and porcine (Miki et al., 1990) membranes yielded single affinity [3 H]-LTB₄ binding sites with K_{DS} of 1.7 nM and 1.0 nM respectively and receptor capacities of 274 fmol mg^{-1} protein and 120 fmol mg^{-1} protein. For mouse spleen membranes the high and low affinity [3H]-LTB4 receptors exhibited binding capacities of 151 fmol mg⁻¹ protein and 354 fmol mg⁻¹ respectively. From our own work and

that of others using human neutrophils we believe the high affinity receptor for LTB₄ to be coupled functionally to chemotaxis and the low affinity receptor for LTB₄ to secretion. Thus the results obtained with mouse spleen membranes suggest that the high affinity [3H]-LTB4 binding site is most probably reflective of a receptor involved in murine neutrophil chemotaxis and the low affinity [3H]-LTB4 receptor involved in secretory phenomena. Previously we have shown the LTB₄ receptor antagonist, CP-105,696 to inhibit [3H]-LTB4 binding to human neutrophil high affinity receptor non-competitively and that was seen for CP-105,696 inhibition of the LTB₄mediated chemotactic response (Showell et al., 1995). As demonstrated here, CP-105,696 is also a non-competitive inhibitor of both [³H]-LTB₄ binding to the high affinity receptor on murine spleen membranes and LTB4-mediated murine neutrophil chemotaxis. Taken together we believe that the findings with CP-105,696 and the above relationship of ligand binding and functional overlap for neutrophil chemotaxis strengthens the argument that the high affinity receptor for LTB₄ is the receptor mediating chemotaxis. With regard to the low affinity receptor for LTB₄ on murine spleen membranes, it appears that this receptor, as was previously shown for human neutrophils (Showell et al., 1995), is coupled to secretory events of murine neutrophils. Again pharmacological evidence using CP-105,696 supports this view. Using murine neutrophils in whole blood, competitive inhibition of LTB4-mediated Mac-1 upregulation was found and occurs at similar concentrations to that for human neutrophil CD11b upregulation in whole blood (Showell et al., 1995). The potency of CP-105,696 in this whole blood assay is underestimated by ~ 2.5 log because of the very high level of protein binding that obtains for this compound (Liston et al., unpublished). In addition [3H]-LTB4 binding to the low affinity LTB₄ receptor on MSM is compe-

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titively inhibited by CP-105,696 as was previously seen for human neutrophils.

Together the data presented here suggest that murine spleen membranes are a relatively easily obtained and useful surrogate for predicting the inhibitory potency and nature of antagonism of prototype LTB₄ antagonists on murine neutrophil functional responses. Furthermore murine and human neutrophils appear to be similarly sensitive to LTB₄ with respect to induction of biological responses and sensitivity to a variety of chemically distinct LTB4 antagonists (Showell et al., unpublished). The value of first being able to demonstrate a concordance between results obtained with human cells and cells from an animal species used for preclinical efficacy testing is considerable. Here we believe we have shown that mice are such a species and evaluation of the LTB₄ antagonist, CP-105,696, in murine models such as collagen-induced arthritis (Griffiths et al., 1995) and experimental allergic encephalomyelitis (Gladue et al., 1995) have provided some compelling data supportive of clinical evaluation of LTB4 antagonists in a variety of human inflammatory diseases.

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