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Pharmacological differences among CysLT₁ receptor antagonists with respect to LTC₄ and LTD₄ in human lung parenchyma

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Abstract

We have previously reported, by means of equilibrium binding studies, the existence of two distinct binding sites with receptor characteristics for LTC₄ and LTD₄ in human lung parenchyma (HLP) membranes using S-decyl-glutathione (S-decyl-GSH) to inhibit LTC₄ binding to a number of non-receptor sites. Recently, we have been able to avoid the use of S-decyl-GSH in kinetic experiments and to characterize a distinctive pharmacological profile for the LTC₄ high affinity binding sites which do not correlate with the ability of both LTD₄ and LTC₄ to contract isolated HLP strips through the CysLT₁ receptor. Here, we report that the most advanced CysLT₁ receptor antagonists, some of which are already in clinical use, displayed a different behavior toward LTC₄ and LTD₄ in HLP. Equilibrium and kinetic binding studies demonstrated the following rank order of potency for ³H-LTD₄ receptor (CysLT₁): zafirlukast = montelukast > LM-1507 = LM-1484 = pranlukast. In addition, LM-1507, LM-1484, pranlukast and montelukast but not zafirlukast are able to interact also with the high affinity site for ³H-LTC₄ (LM-1507 = LM-1484 > pranlukast; montelukast not detectable in the presence of S-decyl-GSH). In this respect, the behavior of the LM antagonists closely resembles that of pranlukast although LM-1507 and LM-1484 display a higher affinity for ³H-LTC₄ sites. Montelukast has an intermediate behavior, inasmuch as its interaction with ³H-LTC₄ sites can be revealed only in kinetic studies, while zafirlukast is totally unable to inhibit ³H-LTC₄ binding. It might be, therefore, most relevant for a complete understanding of the clinical efficacy, besides their nominal potency, of the most advanced CysLT₁ receptor antagonists to consider their pharmacological differences with respect not only to LTD₄/LTE₄, but also to LTC₄. © 2002 Elsevier Science Inc. All rights reserved.

Keywords: Cysteinyl-leukotriene; LM-1507; LM-1484; Montelukast; Human lung parenchyma; Asthma

1. Introduction

Cysteine-containing leukotrienes (Cys-LTs), LTC₄, LTD₄ and LTE₄, originate from the oxidative metabolism of arachidonic acid through a key enzyme, 5-lipoxygenase, in a number of inflammatory cells including eosinophils, basophils, mast cells and macrophages [1,2]. Recently, two different Cys-LT receptor have been cloned, namely CysLT₁ [3,4] and CysLT₂ [5–7], with a distribution that is clearly peculiar for each isoform. In particular, CysLT₁ seems more abundant in HLP, human bronchi and peripheral blood leukocytes and eosinophils [8], while CysLT₂ is more abundant in heart and brain [5]. The identification of the CysLT₁ receptor in the lung is

consistent with the anti-bronchoconstrictive and anti-inflammatory actions of CysLT₁ receptor antagonists.

It is now widely recognized that Cys-LTs play an important role in asthma, participating both to the bronchoconstriction and to the chronic inflammatory component of the disease. Consequently, during the last years the number of structurally different CysLT₁ antagonists has been expanding at an ever increasing rate [9] (Fig. 1) and some are already available for clinical use (zafirlukast, montelukast and pranlukast) or in advance clinical trials (LM-1507, now MEN 91507). In particular, LM-1507 antagonizes in a potent and competitive manner LT-induced contraction in guinea pig ileum and human bronchi *in vitro*, while *in vivo* (administered by the i.v. or oral route) exerts a remarkable and long-lasting antagonism of LTD₄-induced bronchospastic and inflammatory responses.¹ However, all of them, to date, have been

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Abbreviations: LT, leukotriene; Cys-LT, cysteine-containing leukotrienes; S-decyl-GSH, S-decyl-glutathione; HLP, human lung parenchyma; RP-HPLC, reverse phase-high performance liquid chromatography.

¹ Dr. S. Manzini, Personal communications.

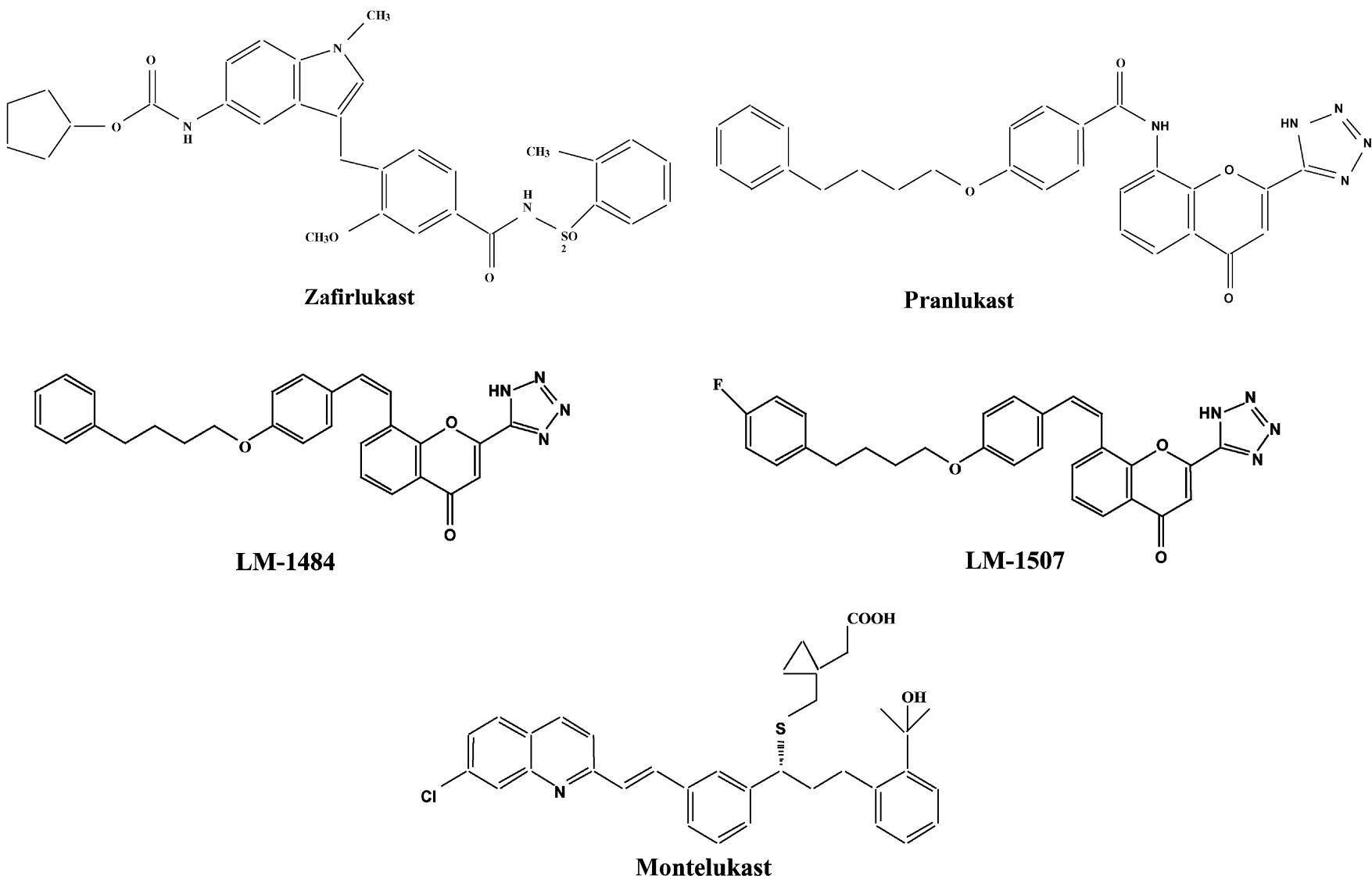


Fig. 1. Chemical structure of CysLT₁ receptor antagonists.

designed to be CysLT₁ receptor antagonists, as it is generally believed that, in humans airways, LTC₄ acts through the same receptor as LTD₄ and LTE₄, namely the CysLT₁ receptor [10–12].

We recently pointed out, however, that in HLP membranes LTC₄ possess a specific high affinity binding site with characteristics distinct from those of LTD₄. In particular, in this tissue two of the classical CysLT₁ antagonists, i.e. zafirlukast and ICI 198,615 (from which zafirlukast has been derived), have different selectivity with respect to ³H-LTC₄ and ³H-LTD₄ at equilibrium, thus suggesting that two different receptors might exist [13]. We also ruled out the hypothesis that this putative LTC₄ receptor might be a CysLT₂. In fact, BAY u9773, the only compound able to recognize both CysLT₁ and CysLT₂ receptors [14], was unable to dissociate ³H-LTC₄ from its high affinity sites [15]. In addition, we have recently characterized a distinctive pharmacological profile for the LTC₄ high affinity binding site which does not correlate with the ability of both LTD₄ and LTC₄ to contract HLP. In fact, in functional experiments all the classical antagonists utilized were able to revert both LTC₄- and LTD₄-induced contractions of isolated HLP strips, thus indicating that both agonist were acting through the same CysLT₁ receptor [15].

It is intriguing to speculate that this putative receptor is implicated in aspects of the asthmatic syndrome different from bronchoconstriction, such as smooth muscle hyperplasia and proliferation or mucus secretion. Indeed, there are data in the literature indicating a proliferative role of Cys-LTs in human airway epithelial [16] or smooth muscle cells [17]. These data not only suggest LTC₄ as a more potent mitogenic stimulus than LTD₄ [16], but also indicate LTD₄ to be a weak agonist with a different pharmacological profile compared to the classical contractile function mediated by the CysLT₁ receptor. In addition, it has been recently reported that an inducible form of CysLT₁ receptor is present in vascular endothelial cells and that aspirin-triggered lipoxin A₄ specifically compete at this receptor site [18]. These data seems to suggest the presence of new sites for action and antagonism in the already complex scenario of inflammatory and anti-inflammatory mediators of asthma.

We think, therefore, that it might be relevant for the complete understanding of the clinical efficacy of the CysLT₁ receptor antagonists to characterize their pharmacological differences with respect not only to LTD₄/LTE₄, but also to LTC₄. We report here that the most advanced CysLT₁ receptor antagonists, which, to date, have been considered a pharmacologically homogeneous class of compounds, actually have a different pharmacological behavior in HLP membranes when challenged against LTC₄ and LTD₄. These data seem to confirm that an “atypical” Cys-LT receptor exists in human airways and that this receptor might possess a different spectrum of activity [17].

2. Materials and methods

2.1. Materials

³H-LTC₄ (164–173 Ci mmol^{−1}) and ³H-LTD₄ (164–173 Ci mmol^{−1}) were purchased from DuPont NEN. LTC₄ and LTD₄ were purchased from Cayman Chemical Co. Zafirlukast (ICI 204,219), pranlukast (ONO 1078), LM-1507 (now MEN 91507) and LM-1484 were from Menarini (LM-1507 is an amphiphilic chromone derivative with no chiral centres, manufactured by a linear synthesis of six steps and developed as a sodium salt); montelukast was from Merck. S-Decyl-GSH, cysteine, glycine, boric acid, serine, Hepes were purchased from Sigma Chemical Co. Filtercount was from Packard Instruments Company. All the reagents used in HPLC analysis were of analytical grade and purchased from Carlo Erba as were GF/C Whatman fiber-glass filters.

2.2. Preparation of HLP membranes

Crude membranes were prepared from macroscopically normal specimens, removed at thoracotomy for lung cancer as previously described [19]. Briefly, specimens were minced and homogenized at 4° in 10 mM Hepes buffer, pH 7.4 (1:24, w/v), centrifuged at 770 g for 10 min and the supernatant centrifuged at 27,000 g for 20 min. The pellet was resuspended, centrifuged under the same condition and finally resuspended in 5% of the homogenization volume. Membrane aliquots were frozen at –80° and stored for no longer than 3 months. Prior to use, serine–borate complex (40 mM, final concentration in the assay, prepared as an equimolar solution of serine and boric acid), cysteine (10 mM) and glycine (10 mM) were added to the membrane suspension to avoid Cys-LT metabolism.

2.3. Reverse phase-high performance liquid chromatography (RP-HPLC)

Labeled and unlabeled leukotriene purity was always assessed by RP-HPLC. Only leukotrienes with a purity grade higher or equal to 95% were utilized. The Beckman HPLC system was equipped with a 110B Solvent Delivery Module, ODS Ultrasphere C18 column and a Programmable Detector Module 166 set at 280 nm. Both labeled and unlabeled leukotrienes were eluted isocratically with a filtered and degassed mixture of CH₃OH:H₂O:CH₃COOH (65:35:0.02 v/v/v), adjusted at pH 5.8 with NH₄OH, at a flow rate of 1 mL min^{−1}. To check the purity of tritiated leukotrienes, fractions were collected every 30 s and the radioactivity profile assessed by liquid scintillation counting (Ultima Gold, Packard).

2.4. Binding studies

Equilibrium binding studies were performed at 25° for 60 min with 0.03–0.5 nM ³H-LTD₄ or 40 min with

0.03–0.5 nM ^3H -LTC₄ and unlabeled homologous or heterologous ligands at the indicated concentrations. A multi-ligand protocol was followed [20,21]. Ten micromolar S-decyl-GSH was present only in the case of ^3H -LTC₄ equilibrium experiments. Time-courses were performed at 25° with 0.5 nM ^3H -LTC₄ or ^3H -LTD₄. Dissociation was induced by adding 1 mM unlabeled leukotriene (homologous dissociation) or 10 μM unlabeled antagonist (heterologous dissociation). In both equilibrium and kinetic studies HLP membranes (0.25 mg per sample), 10 mM HEPES-KOH pH 7.4, 1 mM CaCl₂ and 1 mM MgCl₂ were added to the incubation mixture to achieve a final volume of 250 mL. All the experiments have been performed under control metabolic conditions (see Section 2.2). Unbound ligand was separated by rapid vacuum filtration (Brandel Cell Harvester) onto glass-fiber GF/C filters soaked in 2.5% polyvinylalcohol and the filters were washed twice with 4 mL of HEPES buffer at 4°. Radioactivity was measured in a liquid scintillation counter (Filter Count, Packard).

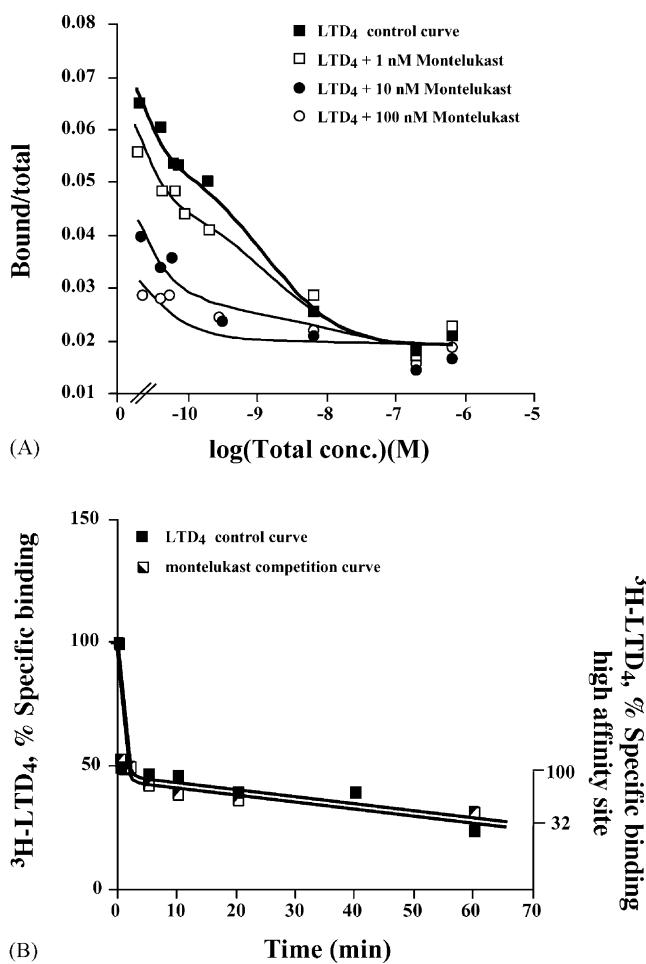


Fig. 2. (A) Equilibrium binding curves for ^3H -LTD₄ in the absence and presence of the indicated concentrations of montelukast (multiligand curves) ($n = 2$). (B) Homologous and heterologous dissociation time-courses. Dissociation of ^3H -LTD₄ was induced by either 1 μM LTD₄ or 10 μM montelukast ($n = 3$). Right axis represents dissociation from the high affinity site only.

2.5. Computer analysis

Analysis of equilibrium ligand binding data was performed by means of the computer program LIGAND [22] that allows simultaneous analysis of replicate experiments. A series of models of increasing complexity involving from one to three binding sites were considered. Analysis of binding data of homologous dissociation time-courses were performed using the program KINFIT II [23], which calculates association and dissociation rates (k_{on} and k_{off}) and the number of binding sites (B_{max}). Binding is expressed as specific bound concentration vs. time. Data from heterologous dissociation time-courses were analyzed using EXPFIT [24], which calculates the coefficients C (amount or percent bound), and R (the apparent rate constant for the specified antagonist). No direct calculation of k_{off} is possible for the heterologous dissociation time-courses. Biphasic dissociation time-courses represent interaction with a heterologous population of sites, where the fast dissociation

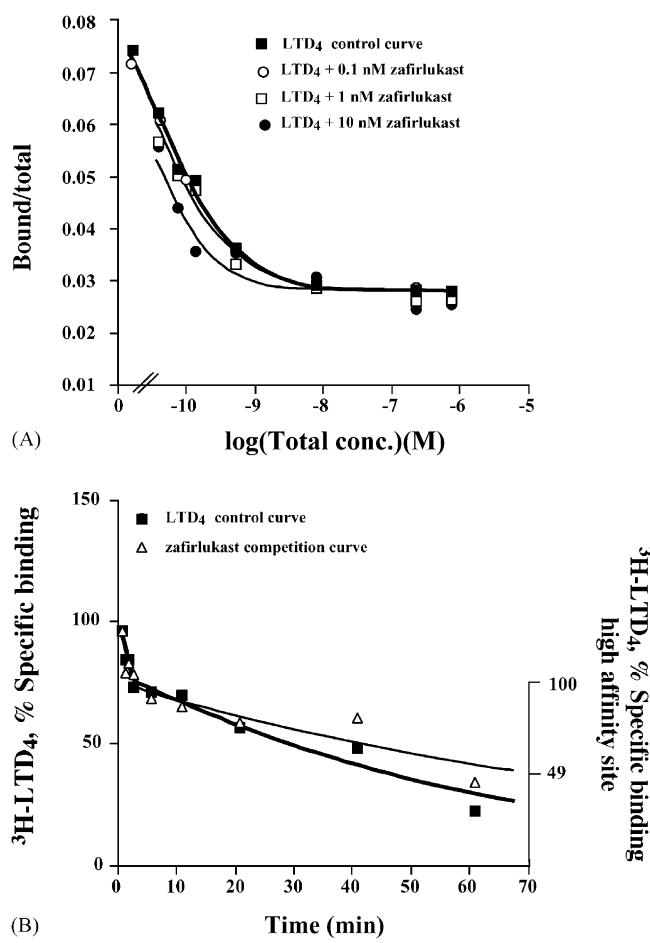


Fig. 3. (A) Equilibrium binding curves for ^3H -LTD₄ in the absence and presence of the indicated concentrations of zafirlukast (multiligand curves) ($n = 2$). (B) Homologous and heterologous dissociation time-courses. Dissociation of ^3H -LTD₄ was induced by either 1 μM LTD₄ or 10 μM zafirlukast ($n = 3$). Right axis represents dissociation from the high affinity site only.

rate represents the low affinity component and the low dissociation rate represents the high affinity component. Antagonist competition is expressed as percentage dissociation specific binding. Different models of increasing complexity were selected using the statistical principle of the “extra sum of squares” [25]. Parameter errors are always expressed in percentage coefficient of variation (%CV). A statistical level of significance of $P < 0.05$ was accepted. All the curves shown were computer generated and represent a typical experiment.

3. Results

3.1. $^3\text{H-LTD}_4$ binding

3.1.1. Equilibrium experiments

Figs. 2A–6A show the multiligand curves (i.e. curves of the labeled ligand in the absence and in the presence of a

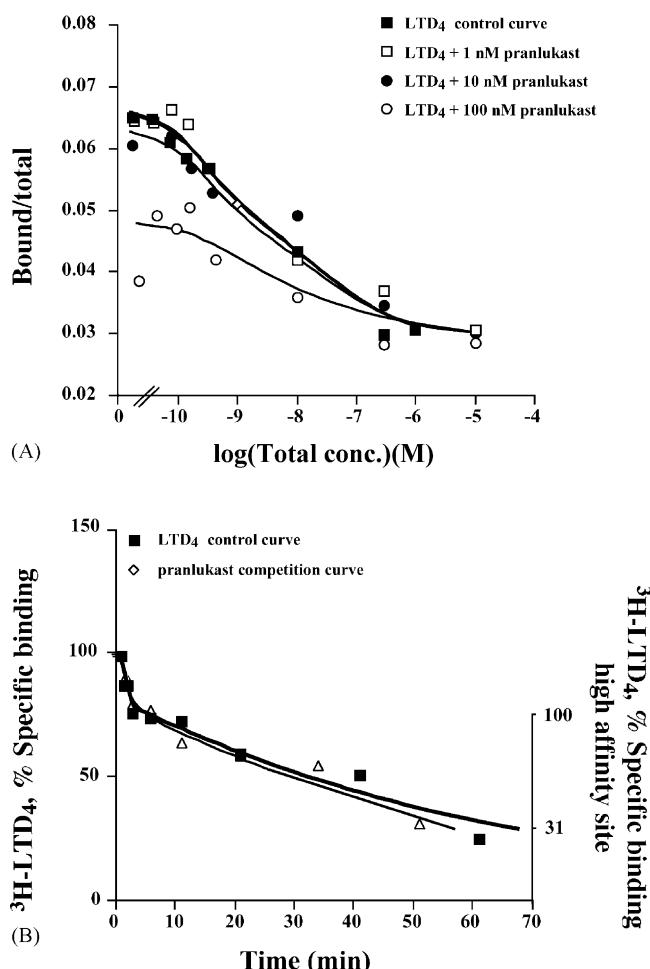


Fig. 4. (A) Equilibrium binding curves for $^3\text{H-LTD}_4$ in the absence and presence of the indicated concentrations of pranlukast (multiligand curves) ($n = 2$). (B) Homologous and heterologous dissociation time-courses. Dissociation of $^3\text{H-LTD}_4$ was induced by either 1 μM LTD₄ or 10 μM pranlukast ($n = 3$). Right axis represents dissociation from the high affinity site only.

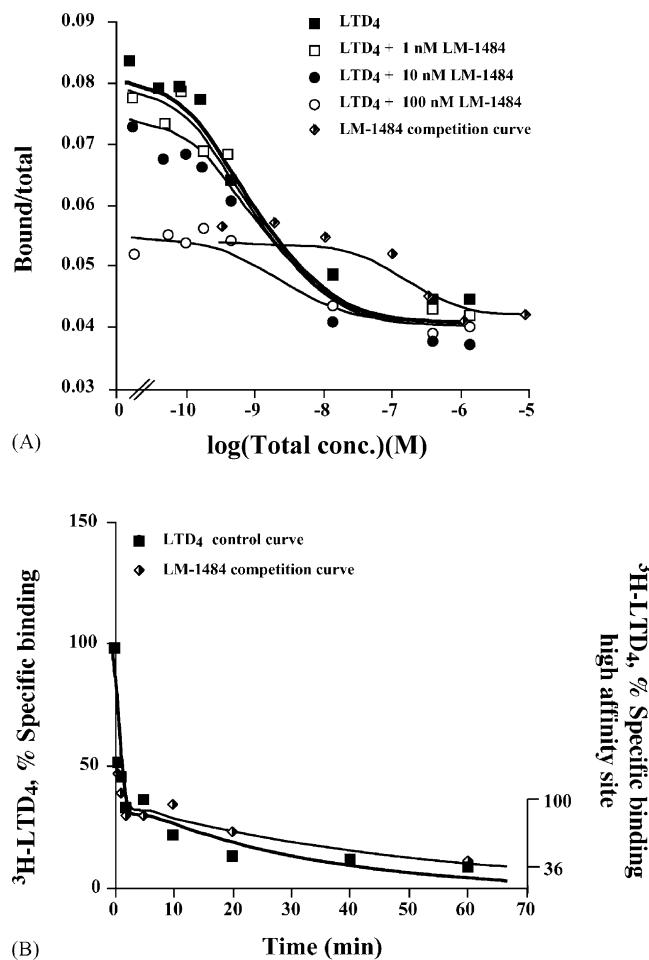


Fig. 5. (A) Equilibrium binding curves for $^3\text{H-LTD}_4$ in the absence and presence of the indicated concentrations of LM-1484 (multiligand curves) ($n = 2$). (B) Homologous and heterologous dissociation time-courses. Dissociation of $^3\text{H-LTD}_4$ was induced by either 1 μM LTD₄ or 10 μM LM-1484 ($n = 3$). Right axis represents dissociation from the high affinity site only.

fixed concentration of the unlabeled ligand) for the CysLT₁ antagonists tested vs. $^3\text{H-LTD}_4$, together with the “mixed curves” for $^3\text{H-LTD}_4$ obtained within the same experiments. As expected, all computerized analysis of such experiments confirmed the presence of two classes of binding sites for $^3\text{H-LTD}_4$ ($K_{d1} = 0.031 \text{ nM} \pm 22\% \text{ CV}$ and $B_{max1} = 0.0034 \text{ pmol mg}^{-1} \text{ protein} \pm 50\% \text{ CV}$; $K_{d2} = 3.8 \text{ nM} \pm 66\% \text{ CV}$ and $B_{max2} = 0.13 \text{ pmol mg}^{-1} \text{ protein} \pm 60\% \text{ CV}$), as predicted by the Ternary Complex Model [26] for agonist binding at a G-protein coupled receptor. On the contrary, all antagonists tested were better fitted by a single class model, again in agreement with Ternary Complex Model. Thus, all CysLT₁ antagonists tested recognize a single binding site in equilibrium studies with the following order of potency: zafirlukast = montelukast > LM-1507 = LM-1484 = pranlukast (montelukast vs. LM-1507, $P < 0.01$). K_i values are reported in Table 1.

3.1.2. Kinetic experiments

In parallel with equilibrium binding studies, we also performed heterologous dissociation time-courses of

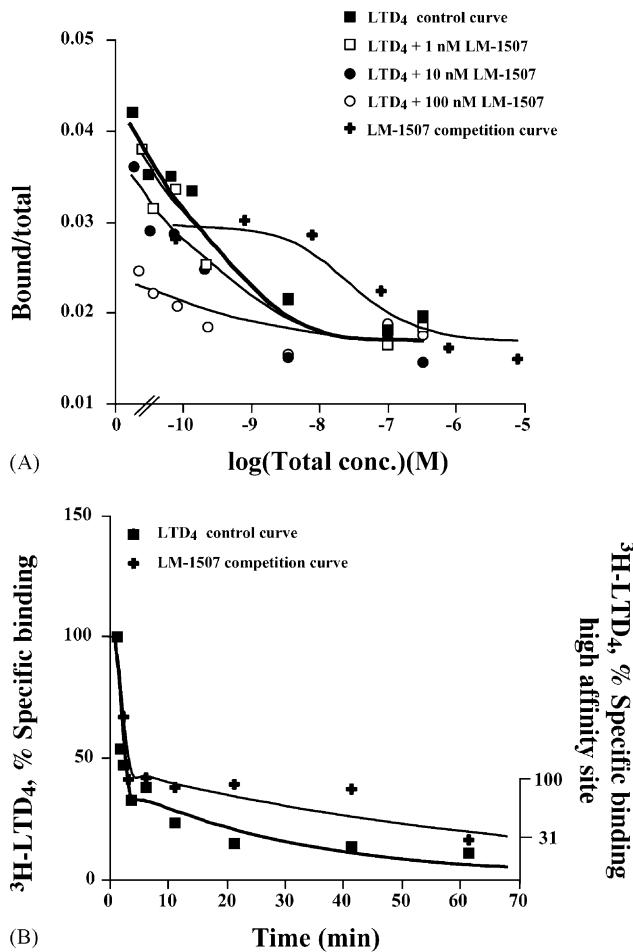


Fig. 6. (A) Equilibrium binding curves for ^3H -LTD₄ in the absence and presence of the indicated concentrations of LM-1507 (multiligand curves) ($n = 2$). (B) Homologous and heterologous dissociation time-courses. Dissociation of ^3H -LTD₄ was induced by either 1 μM LTD₄ or 10 μM LM-1507 ($n = 3$). Right axis represents dissociation from the high affinity site only.

CysLT₁ antagonists in comparison with a ^3H -LTD₄ homologous dissociation curve (Figs. 2B–6B). As expected, all the antagonists tested (Table 1) were able to dissociate ^3H -LTD₄ from its binding sites. In addition, both LTD₄- and antagonist-induced ^3H -LTD₄ dissociation curves were

Table 1
Parameters of the different antagonists for the receptor labeled by ^3H -LTD₄ in human lung parenchyma

Equilibrium experiments $K_{i1} = K_{i2}$ (nM \pm %CV)	Kinetic experiments % dissociation at 60 min	
	High + low affinity sites \pm % CV	
	High affinity site only	
LM-1484	52 \pm 25	88 \pm 7
LM-1507	33 \pm 41	84 \pm 3
Pranlukast	84.7 \pm 23	60 \pm 10
Zafirlukast	1.2 \pm 34	56 \pm 7
Montelukast	2.8 \pm 46	63 \pm 4

Parameters are calculated by means of simultaneous analysis of at least two different experiments.

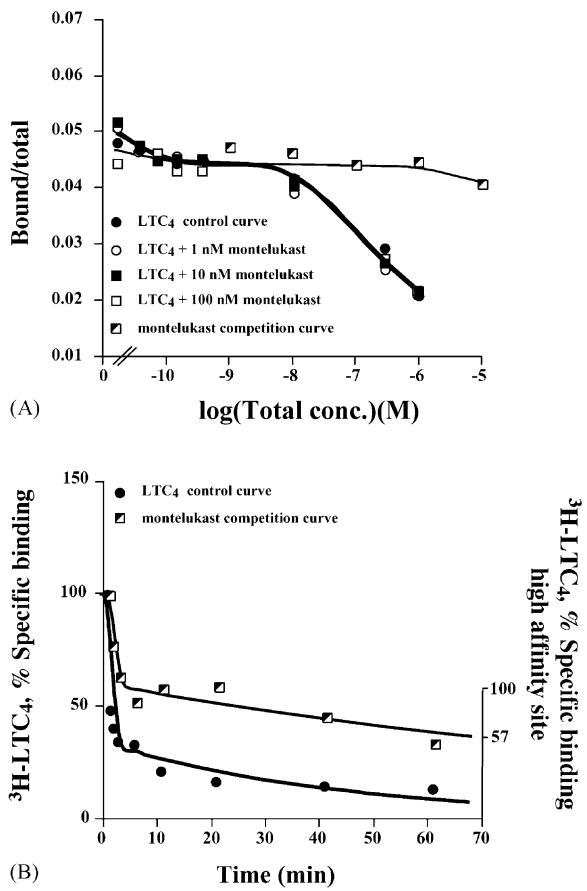


Fig. 7. (A) Equilibrium binding curves for ^3H -LTC₄ in the absence and presence of the indicated concentrations of montelukast (multiligand curves), and classical displacement of ^3H -LTC₄ by montelukast ($n = 3$). (B) Homologous and heterologous dissociation time-courses. Dissociation of ^3H -LTC₄ was induced by either 1 μM LTC₄ or 10 μM montelukast ($n = 3$). Right axis represents dissociation from the high affinity site only.

biphasic ($P < 0.05$), since these experiments represent the dissociation of the labelled ligand (^3H -LTD₄), induced by an unlabelled ligand (LTD₄ or one of the antagonists). The dissociation constant (k_{off1} and k_{off2}) for ^3H -LTD₄ are $1.7 \times 10^{-4} \text{ s}^{-1} \pm 39\% \text{ CV}$ and $4 \times 10^{-3} \text{ s}^{-1} \pm 45\% \text{ CV}$, respectively. On the contrary, no direct calculation of k_{off} was possible for the heterologous dissociation time-courses. Table 1 only reports the percentage dissociation of the high affinity site at 60 min.

3.2. ^3H -LTC₄ binding

3.2.1. Equilibrium experiments

Figs. 7A–11A show the multiligand curves for the CysLT₁ antagonists tested vs. ^3H -LTC₄, together with the “mixed curves” for ^3H -LTC₄ obtained within the same experiments. Computerized analysis of these experiments confirmed the presence of two classes of binding sites for ^3H -LTC₄ ($K_{d1} = 0.21 \text{ nM} \pm 66\% \text{ CV}$ and $B_{max1} = 0.0095 \text{ pmol mg}^{-1} \text{ protein} \pm 50\% \text{ CV}$; $K_{d2} = 113 \text{ nM} \pm 30\% \text{ CV}$ and $B_{max2} = 8.1 \text{ pmol mg}^{-1} \text{ protein} \pm 37\% \text{ CV}$). In equilibrium

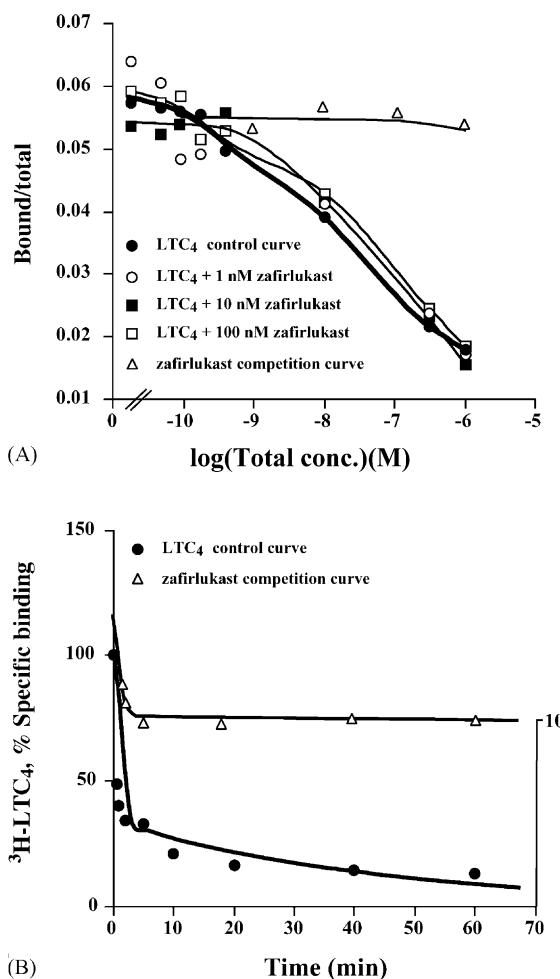


Fig. 8. (A) Equilibrium binding curves for ^3H -LTC₄ in the absence and presence of the indicated concentrations of zafirlukast (multiligand curves), and classical displacement of ^3H -LTC₄ by zafirlukast ($n = 3$). (B) Homologous and heterologous dissociation time-courses. Dissociation of ^3H -LTC₄ was induced by either 1 μM LTC₄ or 10 μM zafirlukast ($n = 3$). Right axis represents dissociation from the high affinity site only.

studies, however, only pranlukast, LM-1484 and LM-1507 (Figs. 9A–11A, respectively) were able to displace ^3H -LTC₄ from its binding sites. These antagonists recognize two classes of binding site, a high affinity one and a low affinity one. The rank order of potency vs. the high affinity class of sites in the presence of S-decyl-GSH is: LM-1507 = LM-1484 > pranlukast (LM-1484 vs. pranlukast, $P < 0.01$). K_i values are reported in Table 2. At equilibrium, zafirlukast was unable to compete with both the sites labeled by LTC₄ (Fig. 8A), while montelukast only marginally interacted with the low affinity one (Fig. 7A).

3.2.2. Kinetic experiments

Dissociation time-courses were performed using ^3H -LTC₄ as labeled ligand. Figs. 7B–11B presents the heterologous dissociation curves for CysLT₁ receptor antagonists vs. ^3H -LTC₄, together with the homologous dissociation curves for LTC₄ obtained within the same experiments. The dissociation constant ($k_{\text{off}1}$ and $k_{\text{off}2}$) for ^3H -LTC₄ are

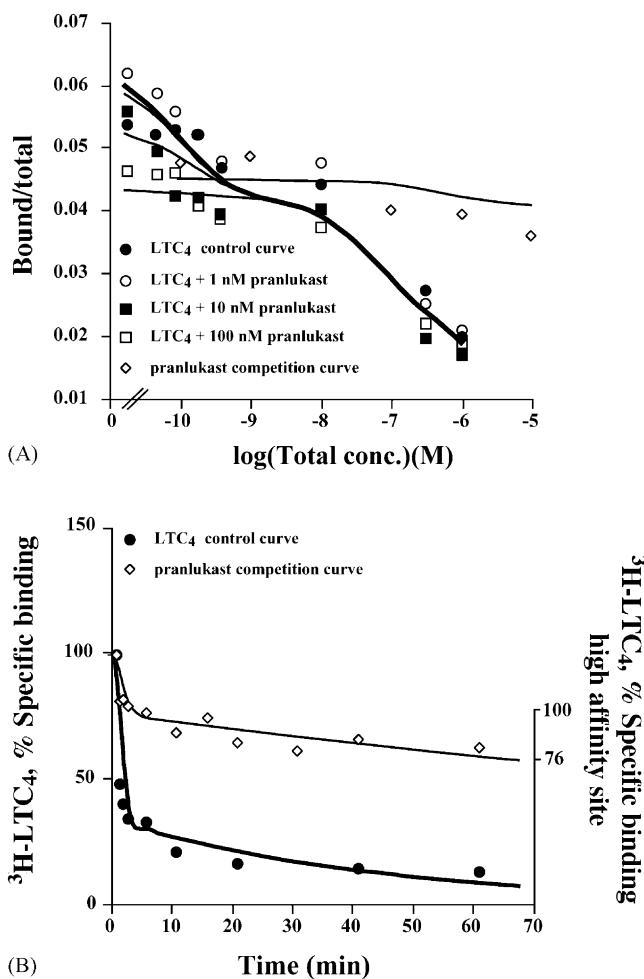


Fig. 9. (A) Equilibrium binding curves for ^3H -LTC₄ in the absence and presence of the indicated concentrations of pranlukast (multiligand curves), and classical displacement of ^3H -LTC₄ by pranlukast ($n = 3$). (B) Homologous and heterologous dissociation time-courses. Dissociation of ^3H -LTC₄ was induced by either 1 μM LTC₄ or 10 μM pranlukast ($n = 3$). Right axis represents dissociation from the high affinity site only.

$7.8 \times 10^{-4} \text{ s}^{-1} \pm 33\% \text{ CV}$ and $2.7 \times 10^{-2} \text{ s}^{-1} \pm 35\% \text{ CV}$, respectively.

LM-1507, LM-1484 and pranlukast displaced ^3H -LTC₄ also in kinetic studies from both high and low affinity sites

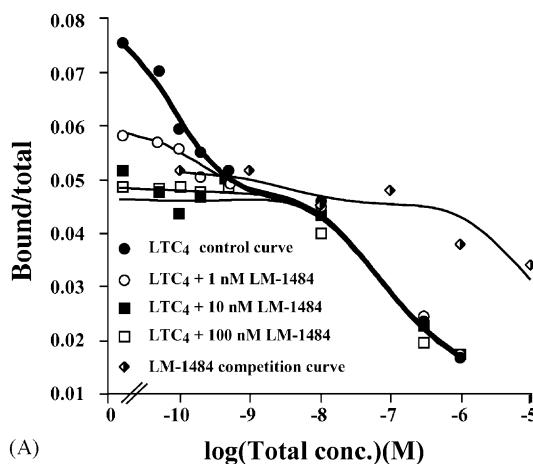
Table 2

Parameters of the different antagonists for the receptor labeled by ^3H -LTC₄ in human lung parenchyma

	Equilibrium experiments		Kinetic experiments
	K_{i1} (nM \pm %CV)	K_{i2} (nM \pm %CV)	% dissociation high affinity site at 60 min \pm % CV
LM-1484	0.55 ± 73	12000 ± 31	40 ± 12
LM-1507	0.26 ± 68	17000 ± 24	38 ± 13
Pranlukast	32 ± 86	27000 ± 95	24 ± 4
Zafirlukast	N.D.	N.D.	N.D.
Montelukast	N.D.	24000 ± 59	43 ± 11

Parameters are calculated by means of simultaneous analysis of at least two different experiments.

N.D. = not detectable.



(A) log(Total conc.) (M)

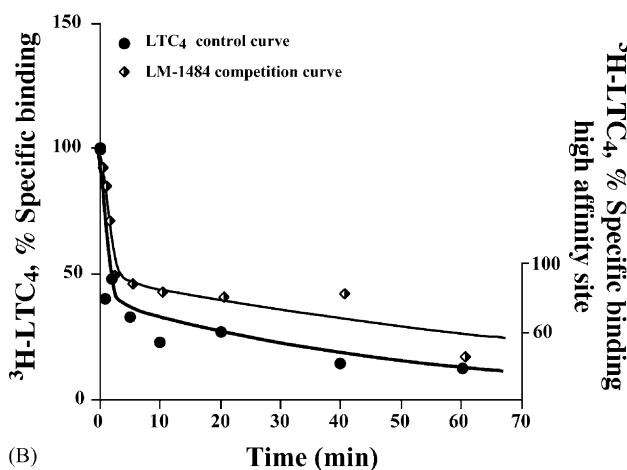
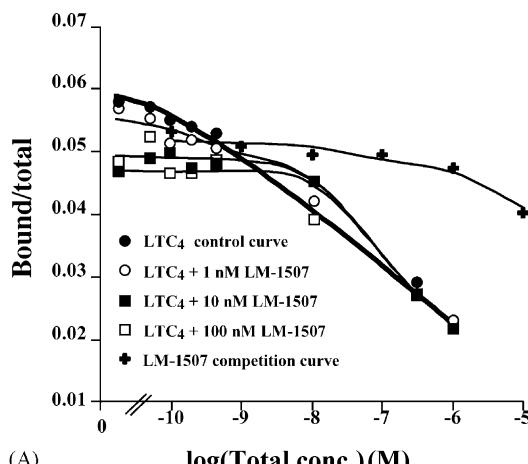


Fig. 10. (A) Equilibrium binding curves for ^3H -LTC₄ in the absence and presence of the indicated concentrations of LM-1484 (multiligand curves), and classical displacement of ^3H -LTC₄ by LM-1484 ($n = 3$). (B) Homologous and heterologous dissociation time-courses. Dissociation of ^3H -LTC₄ was induced by either 1 μM LTC₄ or 10 μM LM-1484 ($n = 3$). Right axis represents dissociation from the high affinity site only.

(Figs. 9B–11B and Table 2), confirming equilibrium data. However, with this experimental protocol also montelukast was able to interact with both sites labeled by ^3H -LTC₄ (Fig. 7B and Table 2). On the contrary, zafirlukast confirmed that it was unable to compete with ^3H -LTC₄ high affinity site, while only a minimal interaction with the low affinity site was detected (Fig. 8B).

4. Discussion

It is well known that LTC₄ predominantly binds to a number of non-receptor sites in cellular membranes [27,28] and, as we previously demonstrated, S-decyl-GSH, a ligand structurally similar to LTC₄ but devoid of either agonist or antagonist activities [29], must be routinely included in ^3H -LTC₄ binding assay at equilibrium to unmask a specific high affinity binding site for LTC₄ [13]. On the contrary, ^3H -LTD₄ binding is basically unaffected by the presence of



(A) log(Total conc.) (M)

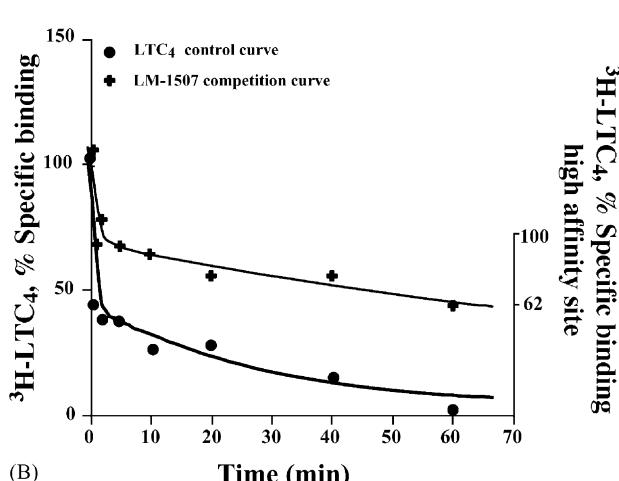


Fig. 11. (A) Equilibrium binding curves for ^3H -LTC₄ in the absence and presence of the indicated concentrations of LM-1507 (multiligand curves), and classical displacement of ^3H -LTC₄ by LM-1507 ($n = 3$). (B) Homologous and heterologous dissociation time-courses. Dissociation of ^3H -LTC₄ was induced by either 1 μM LTC₄ or 10 μM LM-1507 ($n = 3$). Right axis represents dissociation from the high affinity site only.

S-decyl-GSH, as expected for a ligand with low affinity for enzymes such as glutathione-S-transferase. However, we also demonstrated that some antagonists were unable to compete for either ^3H -LTC₄ or ^3H -LTD₄ binding in the presence of S-decyl-GSH, suggesting that this compound interferes with antagonist but not with agonist binding [15]. A possible explanation for these findings could reside in a non total coincidence of agonist and antagonist sites on Cys-LT receptors and in the sterical hindrance of S-decyl-GSH at the antagonist binding site. Alternatively, recently Back *et al.* [30] have hypothesized that this might be due to formation of LTC₄ from LTD₄ in the presence of S-decyl-GSH. Thus, only LTC₄ binding was studied regardless of which agonist was added. This theory substantiates the presence of a different pharmacological profile for LTC₄ and LTD₄ in HLP membranes further supporting our hypothesis of a different LTC₄ receptor in this tissue.

We have also demonstrated that it is possible to avoid the use of S-decyl-GSH by performing heterologous dissoci-

tion time-courses, instead of equilibrium experiments, to study the pharmacological profile of the LTC₄ binding sites [15]. With kinetic experiments, one can perturb the equilibrium with the antagonists to assess their ability to dissociate ³H-LTD₄ or ³H-LTC₄ in the absence of S-decyl-GSH. The only limitation of this type of protocol is that no dissociation constants (k_{off}) or relative potency for the antagonists can be calculated, but only a qualitative analysis can be performed (see Tables 1 and 2). Thus, we have performed the pharmacological characterization of CysLT₁ receptor antagonists by means of both equilibrium and kinetic binding studies.

As expected, all the CysLT₁ receptor antagonists, i.e. zafirlukast, montelukast, LM-1507, LM-1484 and pranlukast, were able to displace ³H-LTD₄ from its binding sites with different potencies both in equilibrium and in kinetic studies. In this respect, zafirlukast and montelukast have the lowest K_i values compared to the others compounds, in agreement with their known clinical potency [31]. Monophasic displacement curves in equilibrium studies suggest a lack of selectivity between the high and low affinity state of CysLT₁ receptor. Indeed, all the antagonists behave as “pure” antagonist with no inverse agonist activity detectable in this assay [32]. Biphasic dissociation time-courses are, however, expected also for nonselective “pure” antagonist, as in heterologous kinetic dissociation curves one actually plot the tracer kinetic (³H-LTD₄, in this case) but not the unlabeled ligand kinetic (the antagonist). For the same reason, the dissociations of ³H-LTD₄ from its high affinity site at a given time are all comparable.

On the contrary, only LM-1507, LM-1484 and pranlukast were able to compete with both ³H-LTC₄ binding sites in equilibrium experiments, even in the presence of S-decyl-GSH. In this respect both the LM compounds have been demonstrated more potent than pranlukast in our experimental conditions. Neither montelukast nor zafirlukast showed any detectable interaction with either site. As mentioned before, the low affinity–high capacity site labeled by ³H-LTC₄ in HLP membranes is of limited interest, inasmuch as it represents a mixture of nonreceptor site and, therefore, K_{i2} values reported in Table 2 have little or no significance. Thus, essentially, both zafirlukast and montelukast are not able to compete with the putative LTC₄ receptor in equilibrium studies. However, kinetic experiments performed in the absence of S-decyl-GSH revealed that montelukast is indeed able to dissociate ³H-LTC₄ also from its high affinity receptor site, therefore, confirming that its inability to do so in equilibrium experiments is most likely due to the interference by S-decyl-GSH. On the contrary, zafirlukast did not show any detectable interaction with either site, confirming its inability to compete with ³H-LTC₄. The kinetic studies confirmed the ability of LM-1507, LM-1484 and pranlukast to interact with both sites labelled by ³H-LTC₄.

A number of CysLT₁ receptor antagonist, i.e. zafirlukast, but also irlukast and the dual CysLT₁–CysLT₂ antagonist

BAY u9773 [15] do not seem to posses any significant antagonism vs. the sites labelled by ³H-LTC₄. Interestingly, only LM-1507 and LM-1484 seem to have a high affinity for the putative LTC₄ receptor, while pranlukast appears to be between 60- and 120-fold less potent, respectively. Indeed, no K_i can be calculated for montelukast, and only functional studies will determine its rank order of potency in comparison to the other antagonist tested. Taken together, these binding data support our hypothesis that classical antagonists should no longer be considered a homogeneous class of compounds with respect to LTC₄ binding sites [15].

Moreover, these results confirm that HLP membranes contain two Cys-LT high-affinity binding sites with different pharmacological profiles. LTD₄ binding sites can be classified as a CysLT₁ receptor, while LTC₄ high affinity binding site is neither a CysLT₁ nor a CysLT₂ receptor. Although a direct evidence to correlate a specific physiological effect with the putative LTC₄ receptor is still lacking, our findings seem to confirm that an “atypical” Cys-LT receptor exists in human airways and that this receptor seems to possess a different spectrum of activity. In particular, it has been shown that LTD₄, besides its known contractile properties, is also able to augment epidermal growth factor-induced human airway smooth muscle cell proliferation. However, this additional effect has a peculiar sensitivity to CysLT₁ receptor antagonist, suggesting that the former phenomenon may be mediated by a Cys-LT receptor distinct from that which mediates LTD₄-induced human airway smooth muscle contraction [17]. It worth notice here that the pharmacological profile of this “atypical” new Cys-LT receptor suggested by these authors is indeed identical to the one proposed by us previously [15] and in this present report for LTC₄.

Many contractile agents [33,34] have been shown to induce proliferation of airway smooth muscle cells in culture and, thus, they might play an important role in stimulating the smooth muscle hypertrophy and hyperplasia associated with asthma. It must be remembered that all Cys-LT receptor antagonists up to date have been developed as specific and selective CysLT₁ receptor antagonists, a characteristic that might demonstrate not particularly convenient. Our finding of antagonist binding to a specific LTC₄ receptor, could contribute to the discovery and development of new and more active drugs with a wider spectrum of action to be used in the treatment of asthma.

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