

## Pharmacological differences among CysLT<sub>1</sub> receptor antagonists with respect to LTC<sub>4</sub> and LTD<sub>4</sub> 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 LTC<sub>4</sub> and LTD<sub>4</sub> in human lung parenchyma (HLP) membranes using *S*-decyl-glutathione (*S*-decyl-GSH) to inhibit LTC<sub>4</sub> 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 LTC<sub>4</sub> high affinity binding sites which do not correlates with the ability of both LTD<sub>4</sub> and LTC<sub>4</sub> to contract isolated HLP strips through the CysLT<sub>1</sub> receptor. Here, we report that the most advanced CysLT<sub>1</sub> receptor antagonists, some of which are already in clinical use, displayed a different behavior toward LTC<sub>4</sub> and LTD<sub>4</sub> in HLP. Equilibrium and kinetic binding studies demonstrated the following rank order of potency for <sup>3</sup>H-LTD<sub>4</sub> receptor (CysLT<sub>1</sub>): 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 <sup>3</sup>H-LTC<sub>4</sub> (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 <sup>3</sup>H-LTC<sub>4</sub> sites. Montelukast has an intermediate behavior, inasmuch as its interaction with <sup>3</sup>H-LTC<sub>4</sub> sites can be revealed only in kinetic studies, while zafirlukast is totally unable to inhibit <sup>3</sup>H-LTC<sub>4</sub> binding. It might be, therefore, most relevant for a complete understanding of the clinical efficacy, besides their nominal potency, of the most advanced CysLT<sub>1</sub> receptor antagonists to consider their pharmacological differences with respect not only to LTD<sub>4</sub>/LTE<sub>4</sub>, but also to LTC<sub>4</sub>. © 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), LTC<sub>4</sub>, LTD<sub>4</sub> and LTE<sub>4</sub>, 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 CysLT<sub>1</sub> [3,4] and CysLT<sub>2</sub> [5–7], with a distribution that is clearly peculiar for each isoform. In particular, CysLT<sub>1</sub> seems more abundant in HLP, human bronchi and peripheral blood leukocytes and eosinophils [8], while CysLT<sub>2</sub> is more abundant in heart and brain [5]. The identification of the CysLT<sub>1</sub> receptor in the lung is

consistent with the anti-bronchoconstrictive and anti-inflammatory actions of CysLT<sub>1</sub> 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 CysLT<sub>1</sub> 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 LTD<sub>4</sub>-induced bronchospastic and inflammatory responses.<sup>1</sup> 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.

<sup>1</sup> Dr. S. Manzini, Personal communications.

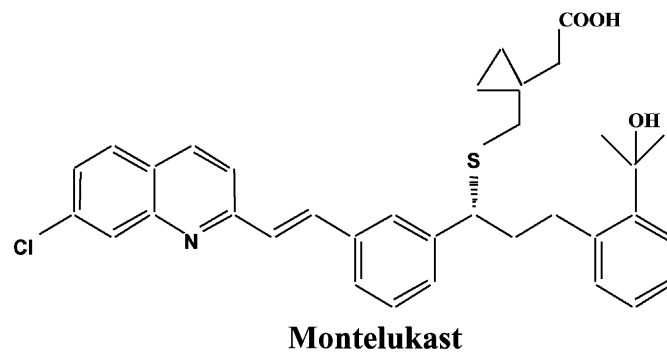
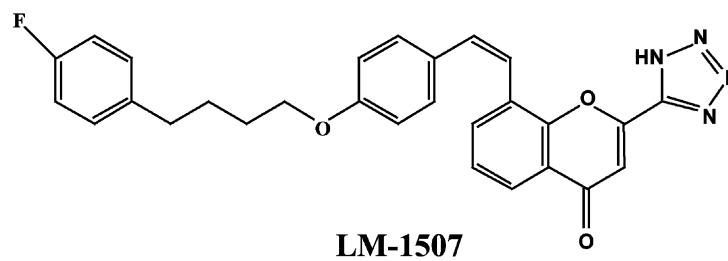
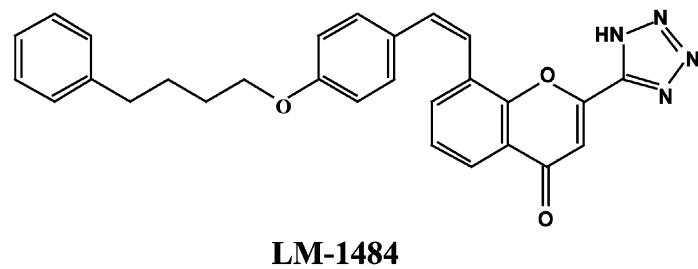
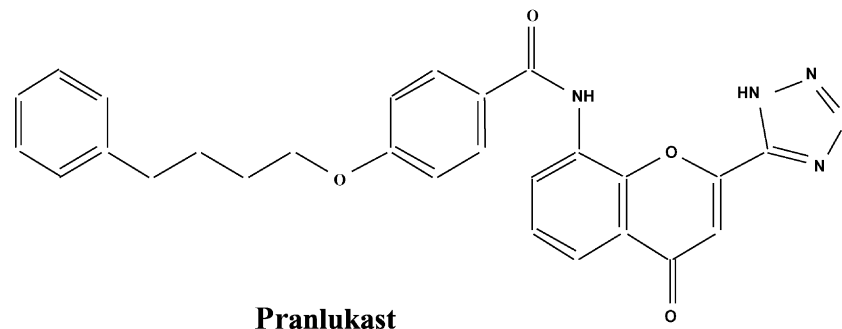
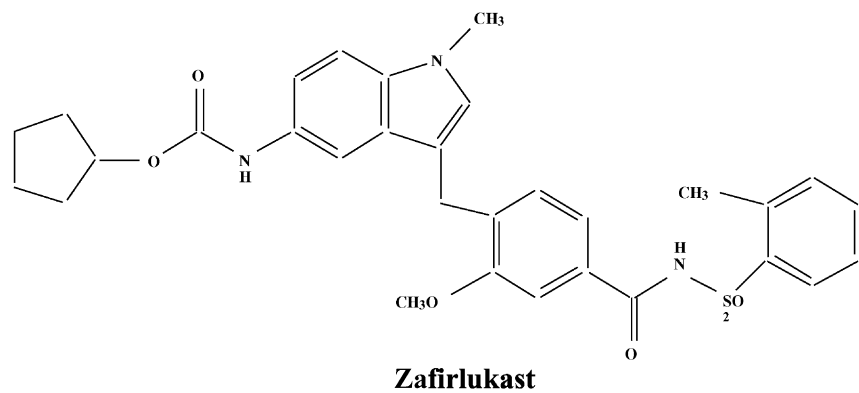


Fig. 1. Chemical structure of CysLT<sub>1</sub> receptor antagonists.

designed to be CysLT<sub>1</sub> receptor antagonists, as it is generally believed that, in humans airways, LTC<sub>4</sub> acts through the same receptor as LTD<sub>4</sub> and LTE<sub>4</sub>, namely the CysLT<sub>1</sub> receptor [10–12].

We recently pointed out, however, that in HLP membranes LTC<sub>4</sub> possess a specific high affinity binding site with characteristics distinct from those of LTD<sub>4</sub>. In particular, in this tissue two of the classical CysLT<sub>1</sub> antagonists, i.e. pobilukast and ICI 198,615 (from which zafirlukast has been derived), have different selectivity with respect to <sup>3</sup>H-LTC<sub>4</sub> and <sup>3</sup>H-LTD<sub>4</sub> at equilibrium, thus suggesting that two different receptors might exist [13]. We also ruled out the hypothesis that this putative LTC<sub>4</sub> receptor might be a CysLT<sub>2</sub>. In fact, BAY u9773, the only compound able to recognize both CysLT<sub>1</sub> and CysLT<sub>2</sub> receptors [14], was unable to dissociate <sup>3</sup>H-LTC<sub>4</sub> from its high affinity sites [15]. In addition, we have recently characterized a distinctive pharmacological profile for the LTC<sub>4</sub> high affinity binding site which does not correlate with the ability of both LTD<sub>4</sub> and LTC<sub>4</sub> to contract HLP. In fact, in functional experiments all the classical antagonists utilized were able to revert both LTC<sub>4</sub>- and LTD<sub>4</sub>-induced contractions of isolated HLP strips, thus indicating that both agonist were acting through the same CysLT<sub>1</sub> 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 LTC<sub>4</sub> as a more potent mitogenic stimulus than LTD<sub>4</sub> [16], but also indicate LTD<sub>4</sub> to be a weak agonist with a different pharmacological profile compared to the classical contractile function mediated by the CysLT<sub>1</sub> receptor. In addition, it has been recently reported that an inducible form of CysLT<sub>1</sub> receptor is present in vascular endothelial cells and that aspirin-triggered lipoxin A<sub>4</sub> 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 CysLT<sub>1</sub> receptor antagonists to characterize their pharmacological differences with respect not only to LTD<sub>4</sub>/LTE<sub>4</sub>, but also to LTC<sub>4</sub>. We report here that the most advanced CysLT<sub>1</sub> 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 LTC<sub>4</sub> and LTD<sub>4</sub>. 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

<sup>3</sup>H-LTC<sub>4</sub> (164–173 Ci mmol<sup>-1</sup>) and <sup>3</sup>H-LTD<sub>4</sub> (164–173 Ci mmol<sup>-1</sup>) were purchased from DuPont NEN. LTC<sub>4</sub> and LTD<sub>4</sub> 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 CH<sub>3</sub>OH:H<sub>2</sub>O:CH<sub>3</sub>COOH (65:35:0.02 v/v/v), adjusted at pH 5.8 with NH<sub>4</sub>OH, at a flow rate of 1 mL min<sup>-1</sup>. 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 <sup>3</sup>H-LTD<sub>4</sub> or 40 min with

0.03–0.5 nM  $^3\text{H-LTC}_4$  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  $^3\text{H-LTC}_4$  equilibrium experiments. Time-courses were performed at 25° with 0.5 nM  $^3\text{H-LTC}_4$  or  $^3\text{H-LTD}_4$ . Dissociation was induced by adding 1 mM unlabeled leukotriene (homologous dissociation) or 10  $\mu\text{M}$  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  $\text{CaCl}_2$  and 1 mM  $\text{MgCl}_2$  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).

## 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 ( $k_{\text{on}}$  and  $k_{\text{off}}$ ) and the number of binding sites ( $B_{\text{max}}$ ). 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  $k_{\text{off}}$  is possible for the heterologous dissociation time-courses. Biphasic dissociation time-courses represent interaction with a heterologous population of sites, where the fast dissociation

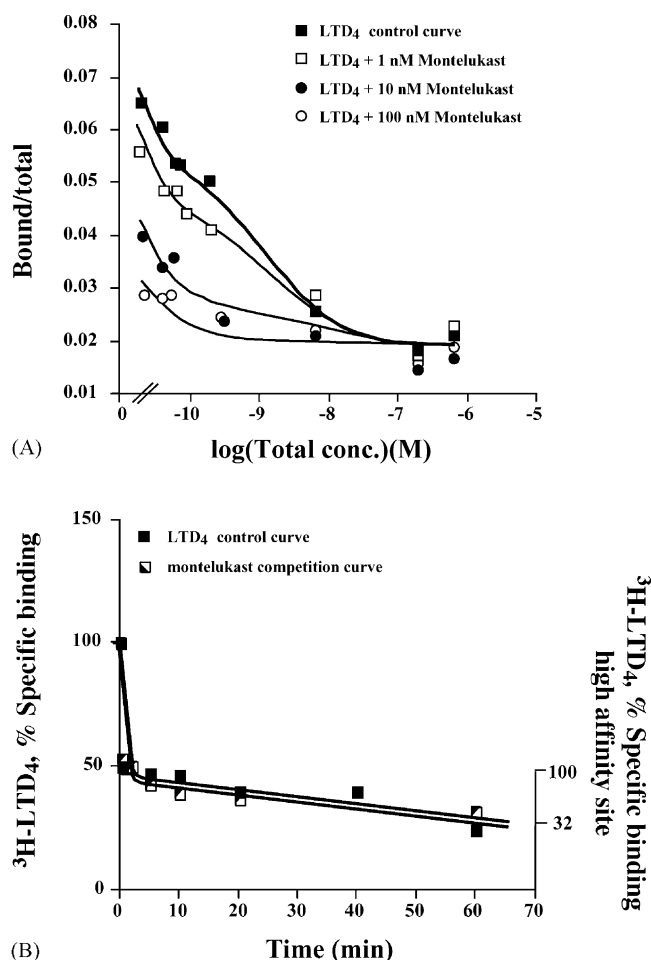


Fig. 2. (A) Equilibrium binding curves for  $^3\text{H-LTD}_4$  in the absence and presence of the indicated concentrations of montelukast (multiligand curves) ( $n = 2$ ). (B) Homologous and heterologous dissociation time-courses. Dissociation of  $^3\text{H-LTD}_4$  was induced by either 1  $\mu\text{M}$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

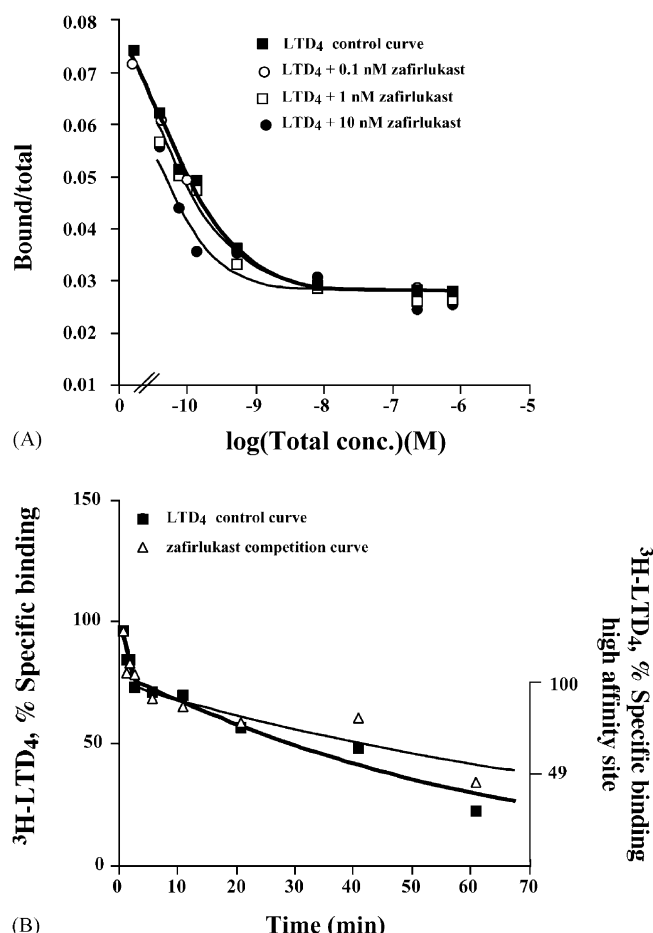


Fig. 3. (A) Equilibrium binding curves for  $^3\text{H-LTD}_4$  in the absence and presence of the indicated concentrations of zafirlukast (multiligand curves) ( $n = 2$ ). (B) Homologous and heterologous dissociation time-courses. Dissociation of  $^3\text{H-LTD}_4$  was induced by either 1  $\mu\text{M}$  LTD<sub>4</sub> or 10  $\mu\text{M}$  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

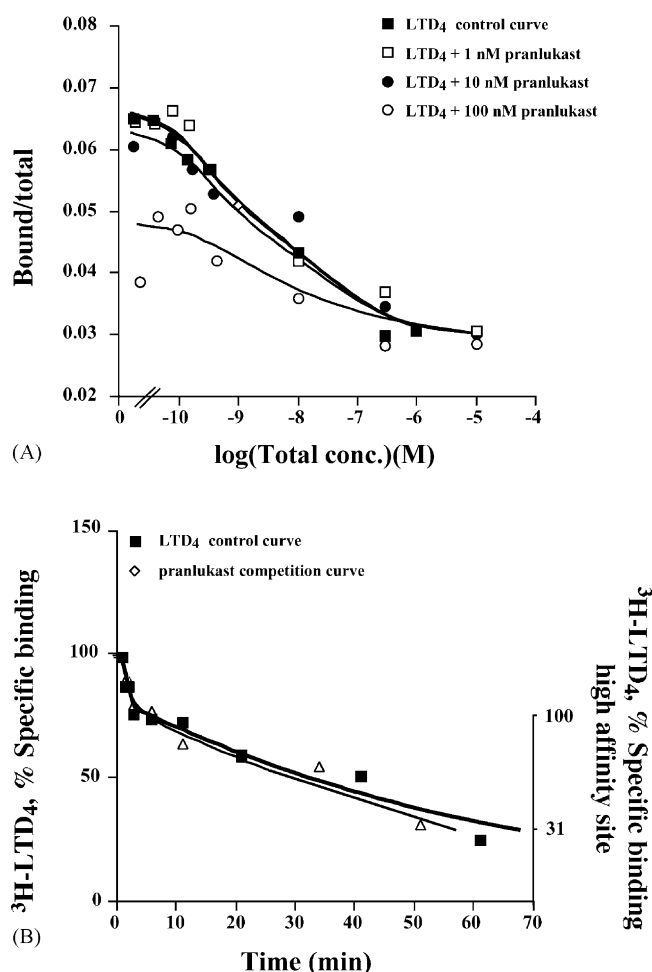
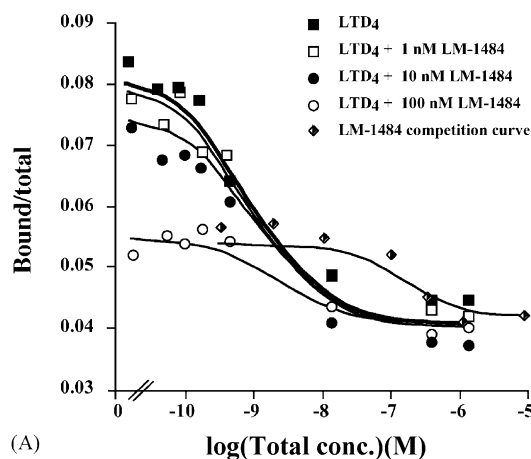
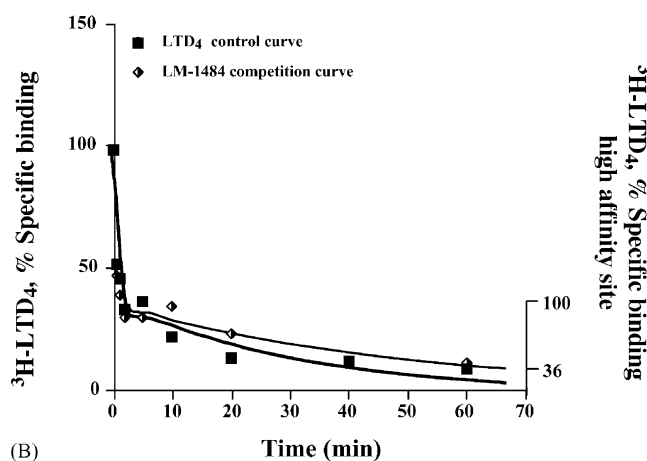


Fig. 4. (A) Equilibrium binding curves for  $^3\text{H-LTD}_4$  in the absence and presence of the indicated concentrations of pranlukast (multiligand curves) ( $n = 2$ ). (B) Homologous and heterologous dissociation time-courses. Dissociation of  $^3\text{H-LTD}_4$  was induced by either 1  $\mu\text{M}$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



(A)



(B)

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  $\mu\text{M}$  LTD $_4$  or 10  $\mu\text{M}$  LM-1484 ( $n = 3$ ). Right axis represents dissociation from the high affinity site only.

fixed concentration of the unlabeled ligand) for the CysLT $_1$  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  $B_{\text{max}1} = 0.0034 \text{ pmol mg}^{-1} \text{ protein} \pm 50\% \text{ CV}$ ;  $K_{d2} = 3.8 \text{ nM} \pm 66\% \text{ CV}$  and  $B_{\text{max}2} = 0.13 \text{ pmol mg}^{-1} \text{ protein} \pm 60\% \text{ CV}$ ), 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 CysLT $_1$  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

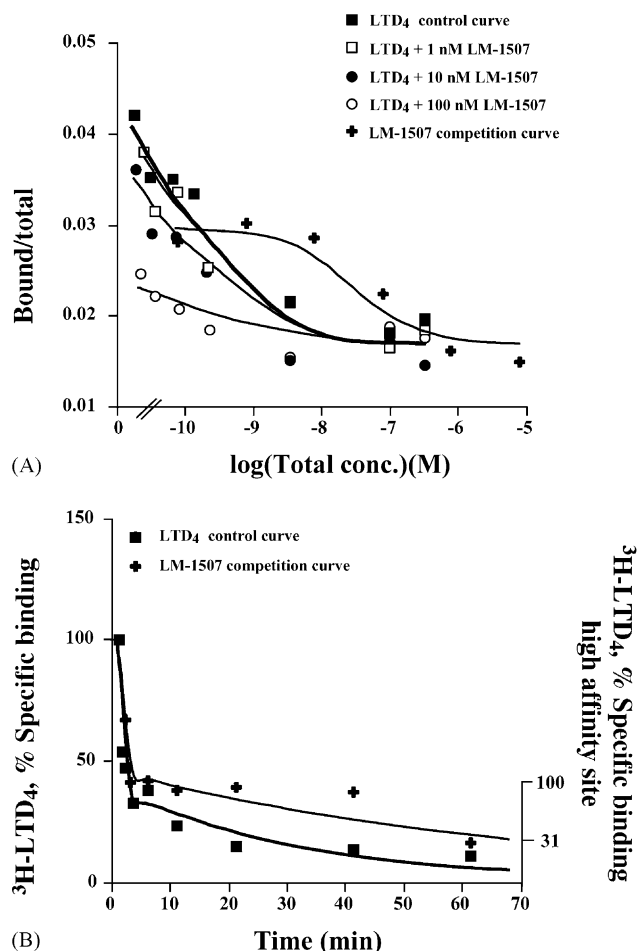


Fig. 6. (A) Equilibrium binding curves for  $^3\text{H-LTD}_4$  in the absence and presence of the indicated concentrations of LM-1507 (multiligand curves) ( $n = 2$ ). (B) Homologous and heterologous dissociation time-courses. Dissociation of  $^3\text{H-LTD}_4$  was induced by either 1  $\mu\text{M}$  LTD $_4$  or 10  $\mu\text{M}$  LM-1507 ( $n = 3$ ). Right axis represents dissociation from the high affinity site only.

CysLT $_1$  antagonists in comparison with a  $^3\text{H-LTD}_4$  homologous dissociation curve (Figs. 2B–6B). As expected, all the antagonists tested (Table 1) were able to dissociate  $^3\text{H-LTD}_4$  from its binding sites. In addition, both LTD $_4$ - and antagonist-induced  $^3\text{H-LTD}_4$  dissociation curves were

Table 1

Parameters of the different antagonists for the receptor labeled by  $^3\text{H-LTD}_4$  in human lung parenchyma

	Equilibrium experiments $K_{d1} = K_{d2}$ (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	64
LM-1507	33 $\pm$ 41	84 $\pm$ 3	69
Pranlukast	84.7 $\pm$ 23	60 $\pm$ 10	69
Zafirlukast	1.2 $\pm$ 34	56 $\pm$ 7	51
Montelukast	2.8 $\pm$ 46	63 $\pm$ 4	68

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

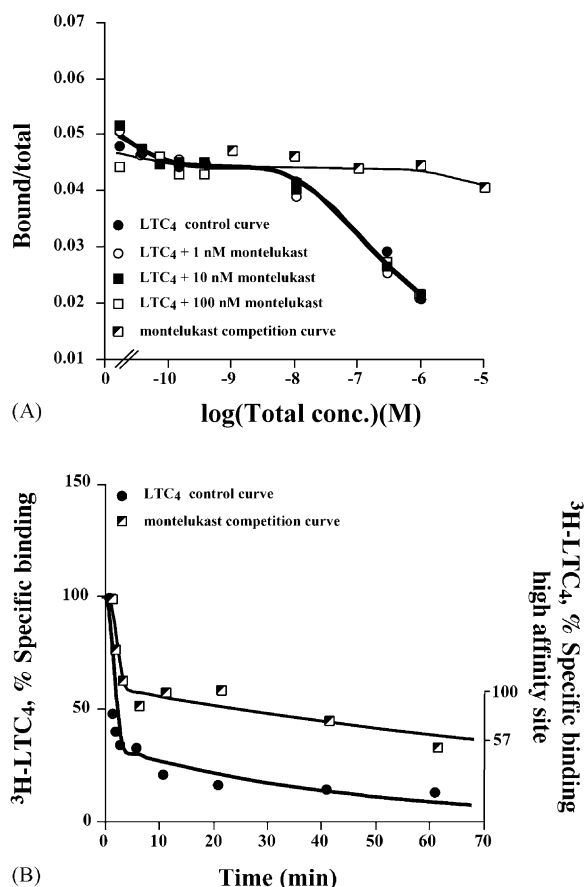


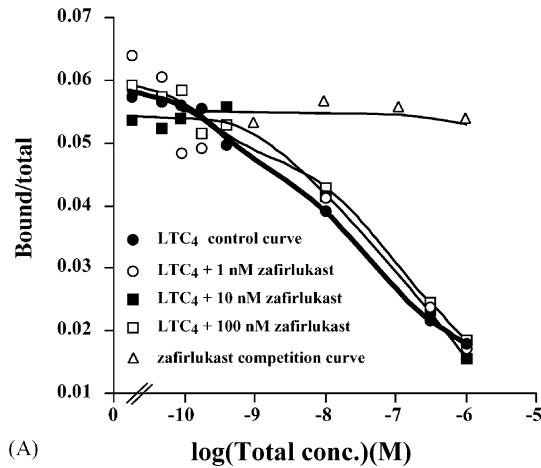
Fig. 7. (A) Equilibrium binding curves for  $^3\text{H-LTC}_4$  in the absence and presence of the indicated concentrations of montelukast (multiligand curves), and classical displacement of  $^3\text{H-LTC}_4$  by montelukast ( $n = 3$ ). (B) Homologous and heterologous dissociation time-courses. Dissociation of  $^3\text{H-LTC}_4$  was induced by either 1  $\mu\text{M}$  LTC $_4$  or 10  $\mu\text{M}$  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 ( $^3\text{H-LTD}_4$ ), induced by an unlabelled ligand (LTD $_4$  or one of the antagonists). The dissociation constant ( $k_{\text{off}1}$  and  $k_{\text{off}2}$ ) for  $^3\text{H-LTD}_4$  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  $k_{\text{off}}$  was possible for the heterologous dissociation time-courses. Table 1 only reports the percentage dissociation of the high affinity site at 60 min.

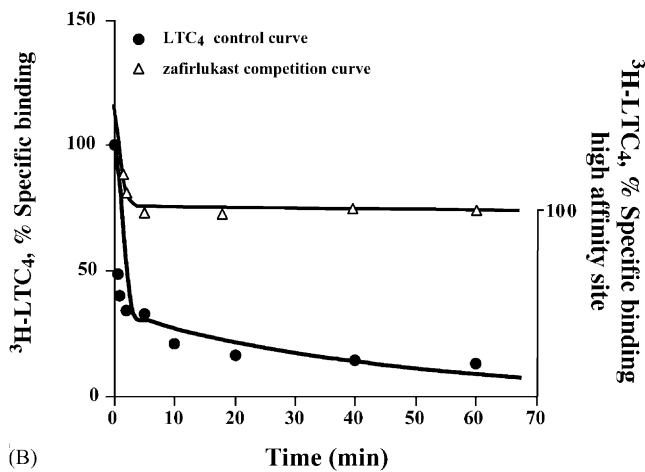
### 3.2. $^3\text{H-LTC}_4$ binding

#### 3.2.1. Equilibrium experiments

Figs. 7A–11A show the multiligand curves for the CysLT $_1$  antagonists tested vs.  $^3\text{H-LTC}_4$ , together with the “mixed curves” for  $^3\text{H-LTC}_4$  obtained within the same experiments. Computerized analysis of these experiments confirmed the presence of two classes of binding sites for  $^3\text{H-LTC}_4$  ( $K_{d1} = 0.21 \text{ nM} \pm 66\% \text{ CV}$  and  $B_{\text{max}1} = 0.0095 \text{ pmol mg}^{-1} \text{ protein} \pm 50\% \text{ CV}$ ;  $K_{d2} = 113 \text{ nM} \pm 30\% \text{ CV}$  and  $B_{\text{max}2} = 8.1 \text{ pmol mg}^{-1} \text{ protein} \pm 37\% \text{ CV}$ ). In equilibrium



(A)



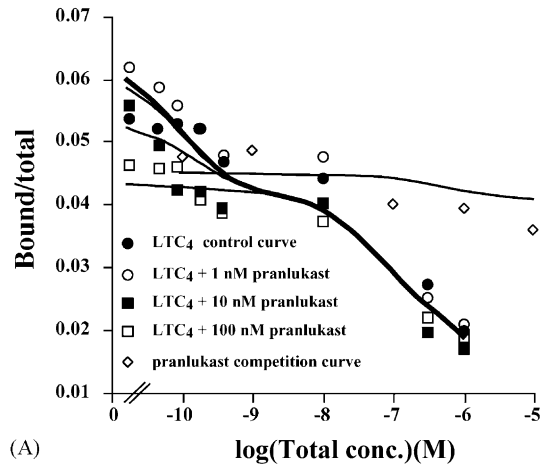
(B)

Fig. 8. (A) Equilibrium binding curves for  $^3\text{H-LTC}_4$  in the absence and presence of the indicated concentrations of zafirlukast (multiligand curves), and classical displacement of  $^3\text{H-LTC}_4$  by zafirlukast ( $n = 3$ ). (B) Homologous and heterologous dissociation time-courses. Dissociation of  $^3\text{H-LTC}_4$  was induced by either  $1 \mu\text{M LTC}_4$  or  $10 \mu\text{M zafirlukast}$  ( $n = 3$ ). Right axis represents dissociation from the high affinity site only.

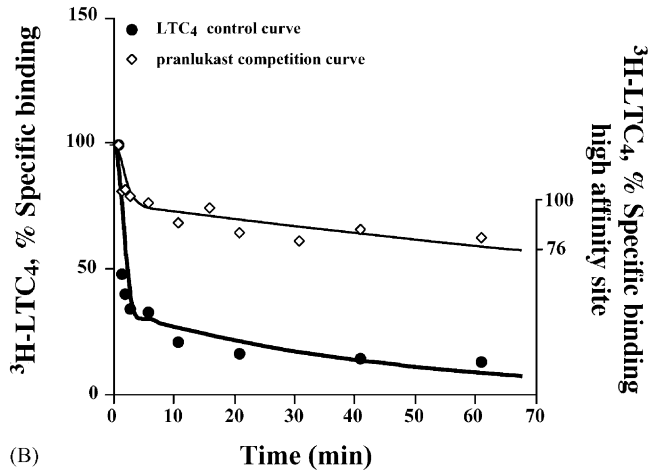
studies, however, only pranlukast, LM-1484 and LM-1507 (Figs. 9A–11A, respectively) were able to displace  $^3\text{H-LTC}_4$  from its binding sites. These antagonists recognize two classes of binding site, a high affinity one and a low affinity one. The rank order of potency vs. the high affinity class 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  $\text{LTC}_4$  (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  $^3\text{H-LTC}_4$  as labeled ligand. Figs. 7B–11B presents the heterologous dissociation curves for  $\text{CysLT}_1$  receptor antagonists vs.  $^3\text{H-LTC}_4$ , together with the homologous dissociation curves for  $\text{LTC}_4$  obtained within the same experiments. The dissociation constant ( $k_{\text{off}1}$  and  $k_{\text{off}2}$ ) for  $^3\text{H-LTC}_4$  are



(A)



(B)

Fig. 9. (A) Equilibrium binding curves for  $^3\text{H-LTC}_4$  in the absence and presence of the indicated concentrations of pranlukast (multiligand curves), and classical displacement of  $^3\text{H-LTC}_4$  by pranlukast ( $n = 3$ ). (B) Homologous and heterologous dissociation time-courses. Dissociation of  $^3\text{H-LTC}_4$  was induced by either  $1 \mu\text{M LTC}_4$  or  $10 \mu\text{M pranlukast}$  ( $n = 3$ ). 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  $^3\text{H-LTC}_4$  also in kinetic studies from both high and low affinity sites

Table 2

Parameters of the different antagonists for the receptor labeled by  $^3\text{H-LTC}_4$  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	$0.55 \pm 73$	$12000 \pm 31$	$40 \pm 12$
LM-1507	$0.26 \pm 68$	$17000 \pm 24$	$38 \pm 13$
Pranlukast	$32 \pm 86$	$27000 \pm 95$	$24 \pm 4$
Zafirlukast	N.D.	N.D.	N.D.
Montelukast	N.D.	$24000 \pm 59$	$43 \pm 11$

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

N.D. = not detectable.

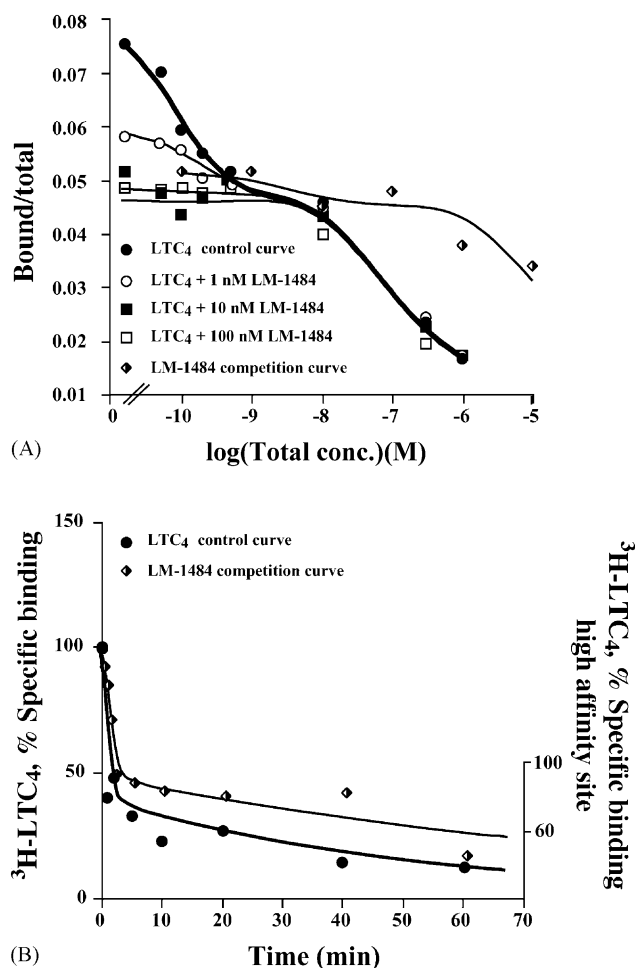


Fig. 10. (A) Equilibrium binding curves for <sup>3</sup>H-LTC<sub>4</sub> in the absence and presence of the indicated concentrations of LM-1484 (multiligand curves), and classical displacement of <sup>3</sup>H-LTC<sub>4</sub> by LM-1484 ( $n = 3$ ). (B) Homologous and heterologous dissociation time-courses. Dissociation of <sup>3</sup>H-LTC<sub>4</sub> was induced by either 1  $\mu$ M LTC<sub>4</sub> or 10  $\mu$ M 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 <sup>3</sup>H-LTC<sub>4</sub> (Fig. 7B and Table 2). On the contrary, zafirlukast confirmed that it was unable to compete with <sup>3</sup>H-LTC<sub>4</sub> high affinity site, while only a minimal interaction with the low affinity site was detected (Fig. 8B).

#### 4. Discussion

It is well known that LTC<sub>4</sub> 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 LTC<sub>4</sub> but devoid of either agonist or antagonist activities [29], must be routinely included in <sup>3</sup>H-LTC<sub>4</sub> binding assay at equilibrium to unmask a specific high affinity binding site for LTC<sub>4</sub> [13]. On the contrary, <sup>3</sup>H-LTD<sub>4</sub> binding is basically unaffected by the presence of

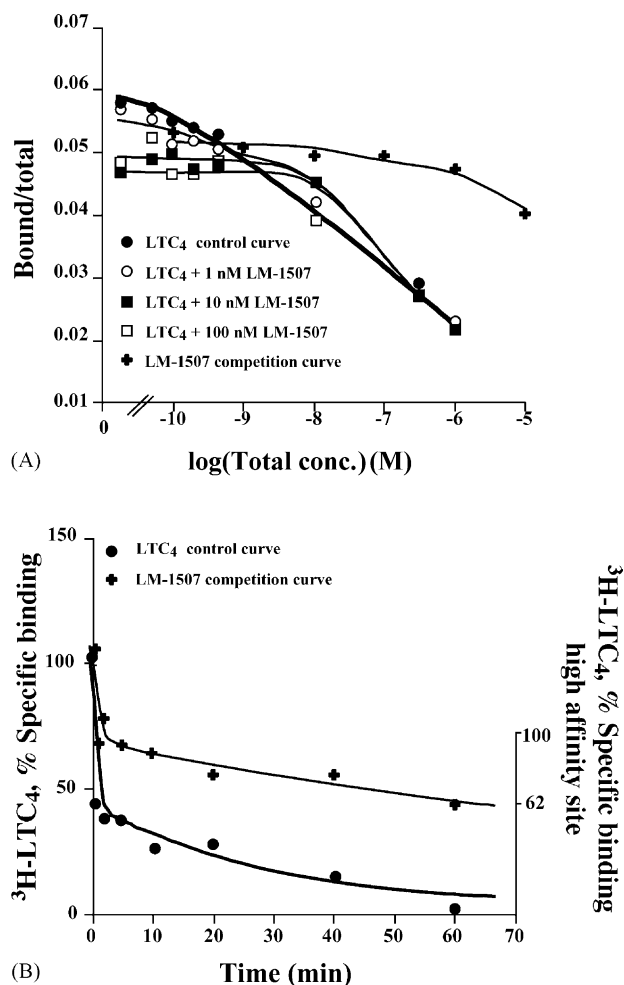


Fig. 11. (A) Equilibrium binding curves for <sup>3</sup>H-LTC<sub>4</sub> in the absence and presence of the indicated concentrations of LM-1507 (multiligand curves), and classical displacement of <sup>3</sup>H-LTC<sub>4</sub> by LM-1507 ( $n = 3$ ). (B) Homologous and heterologous dissociation time-courses. Dissociation of <sup>3</sup>H-LTC<sub>4</sub> was induced by either 1  $\mu$ M LTC<sub>4</sub> or 10  $\mu$ M 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 <sup>3</sup>H-LTC<sub>4</sub> or <sup>3</sup>H-LTD<sub>4</sub> 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 LTC<sub>4</sub> from LTD<sub>4</sub> in the presence of *S*-decyl-GSH. Thus, only LTC<sub>4</sub> binding was studied regardless of which agonist was added. This theory substantiates the presence of a different pharmacological profile for LTC<sub>4</sub> and LTD<sub>4</sub> in HLP membranes further supporting our hypothesis of a different LTC<sub>4</sub> receptor in this tissue.

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



tion time-courses, instead of equilibrium experiments, to study the pharmacological profile of the LTC<sub>4</sub> binding sites [15]. With kinetic experiments, one can perturb the equilibrium with the antagonists to assess their ability to dissociate <sup>3</sup>H-LTD<sub>4</sub> or <sup>3</sup>H-LTC<sub>4</sub> in the absence of *S*-decyl-GSH. The only limitation of this type of protocol is that no dissociation constants (*k*<sub>off</sub>) 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 CysLT<sub>1</sub> receptor antagonists by means of both equilibrium and kinetic binding studies.

As expected, all the CysLT<sub>1</sub> receptor antagonists, i.e. zafirlukast, montelukast, LM-1507, LM-1484 and pranlukast, were able to displace <sup>3</sup>H-LTD<sub>4</sub> from its binding sites with different potencies both in equilibrium and in kinetic studies. In this respect, zafirlukast and montelukast have the lowest *K*<sub>i</sub> 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 CysLT<sub>1</sub> 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 (<sup>3</sup>H-LTD<sub>4</sub>, in this case) but not the unlabeled ligand kinetic (the antagonist). For the same reason, the dissociations of <sup>3</sup>H-LTD<sub>4</sub> 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 <sup>3</sup>H-LTC<sub>4</sub> 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 <sup>3</sup>H-LTC<sub>4</sub> in HLP membranes is of limited interest, inasmuch as it represents a mixture of nonreceptor site and, therefore, *K*<sub>i2</sub> values reported in Table 2 have little or no significance. Thus, essentially, both zafirlukast and montelukast are not able to compete with the putative LTC<sub>4</sub> receptor in equilibrium studies. However, kinetic experiments performed in the absence of *S*-decyl-GSH revealed that montelukast is indeed able to dissociate <sup>3</sup>H-LTC<sub>4</sub> 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 <sup>3</sup>H-LTC<sub>4</sub>. The kinetic studies confirmed the ability of LM-1507, LM-1484 and pranlukast to interact with both sites labelled by <sup>3</sup>H-LTC<sub>4</sub>.

A number of CysLT<sub>1</sub> receptor antagonist, i.e. zafirlukast, but also iralukast and the dual CysLT<sub>1</sub>–CysLT<sub>2</sub> antagonist

BAY u9773 [15] do not seem to possess any significant antagonism vs. the sites labelled by <sup>3</sup>H-LTC<sub>4</sub>. Interestingly, only LM-1507 and LM-1484 seem to have a high affinity for the putative LTC<sub>4</sub> receptor, while pranlukast appears to be between 60- and 120-fold less potent, respectively. Indeed, no *K*<sub>i</sub> 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 LTC<sub>4</sub> binding sites [15].

Moreover, these results confirm that HLP membranes contain two Cys-LT high-affinity binding sites with different pharmacological profiles. LTD<sub>4</sub> binding sites can be classified as a CysLT<sub>1</sub> receptor, while LTC<sub>4</sub> high affinity binding site is neither a CysLT<sub>1</sub> nor a CysLT<sub>2</sub> receptor. Although a direct evidence to correlate a specific physiological effect with the putative LTC<sub>4</sub> 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 LTD<sub>4</sub>, 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 CysLT<sub>1</sub> receptor antagonist, suggesting that the former phenomenon may be mediated by a Cys-LT receptor distinct from that which mediates LTD<sub>4</sub>-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 LTC<sub>4</sub>.

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 CysLT<sub>1</sub> receptor antagonists, a characteristic that might demonstrate not particularly convenient. Our finding of antagonist binding to a specific LTC<sub>4</sub> 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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