

DIARYL ETHER/CARBOXYLIC ACID DERIVATIVES OF LY255283: RECEPTOR ANTAGONISTS OF LEUKOTRIENE B₄

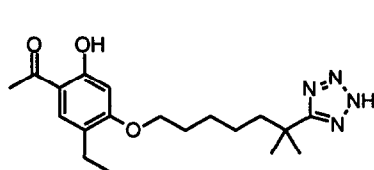
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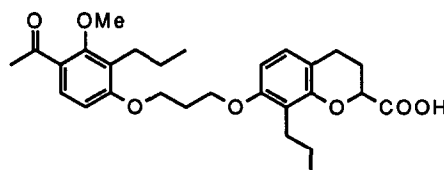
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Abstract: The preparation and activity of a series of hydroxyacetophenone/diaryl ether LTB₄ receptor antagonists are described. The key acid-substituted diaryl ether moiety is discussed in relation to the spatial and functional group requirements of the LTB₄ receptor.

There is increasing evidence that the pro-inflammatory lipid LTB₄ may play an important, receptor-mediated, role in disease states ranging from asthma to inflammatory bowel disease (IBD).¹ The development of potent and specific antagonists² against LTB₄ is critical to elucidating its function in the initiation of inflammatory pathogenesis. Two recently developed LTB₄ receptor antagonists, LY255283³ and SC-41930,⁴ are hydroxyacetophenone-based



LY255283

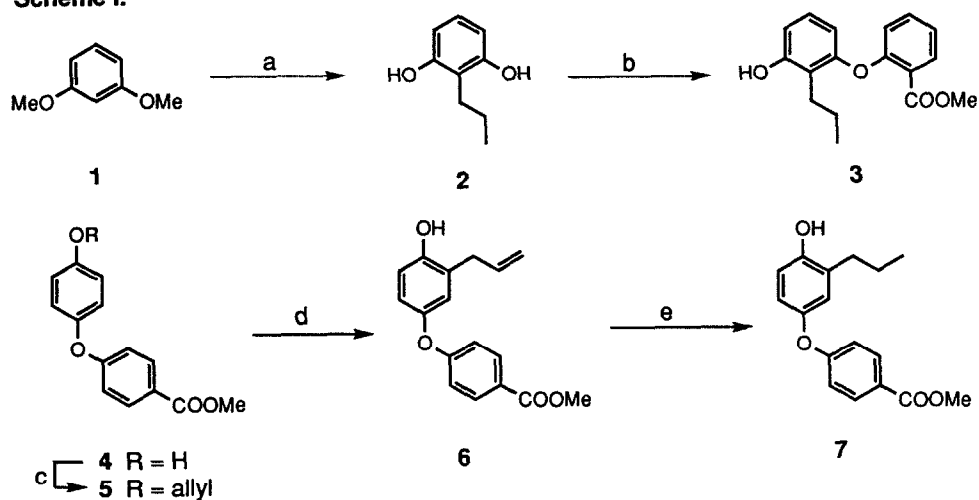


SC-41930

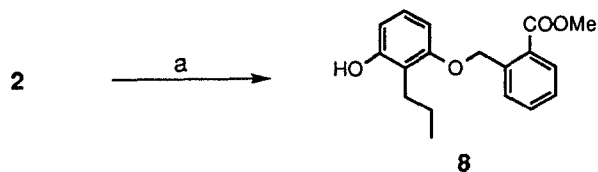
structures that evolved from two previous series of LTD₄/LTE₄ antagonists. We became interested in synthesizing new acid components for LY255283 with a goal of improving binding affinity for the LTB₄ receptor. Examination of the acid-bearing scaffolding of SC41930 revealed a constrained structure in which the carboxyl group is directed out-of-plane relative to a chroman aromatic ring containing a secondary lipophilic group.⁵ We reasoned that an acid-substituted diaryl ether moiety, similar to the chroman unit of SC-41930, would allow sufficient flexibility to enhance receptor binding. Accordingly, a short series of such diaryl ether units were synthesized and merged into the LY255283 hydroxyacetophenone structure to create a new series of antagonists.

Schemes I-III are representative of the chemistry used to prepare the target compounds. A classical Ullmann procedure^{2j} was used to construct the diaryl ethers as shown for phenol 3 (Scheme I) in modest yields.⁶ Appendage of an *n*-propyl side chain was accomplished either through direct alkylation (conversion of 1 to 2), or Claisen rearrangement and reduction beginning with an allyloxy derivative (conversion of 5 to 7).^{2d} Phenol 8, the appropriate acid

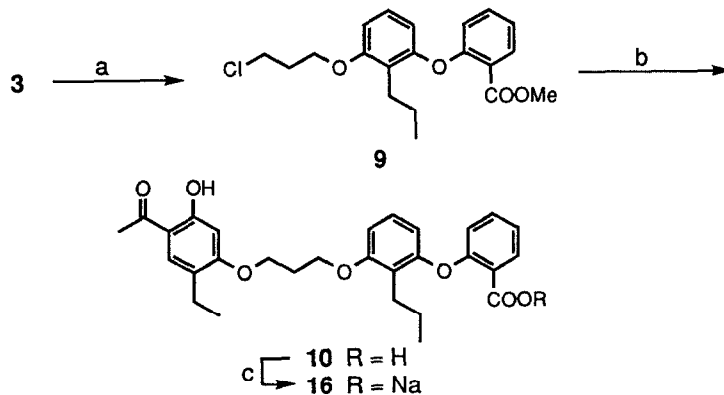
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Scheme I.

a) (i) n -BuLi/ n -PrI/THF (90%), (ii) py-HCl/180 °C (96%); b) methyl 2-iodobenzoate/ $\text{Cu}^0/\text{K}_2\text{CO}_3$ /py/reflux (36%); c) allyl iodide/ K_2CO_3 /2-butanone/DMSO (78%); d) neat/225 °C (83%); e) H_2 /10% Pd(C)/MeOH (95%).

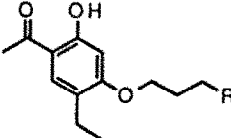
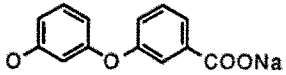
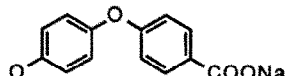
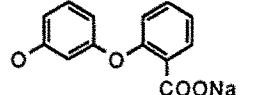
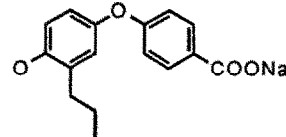
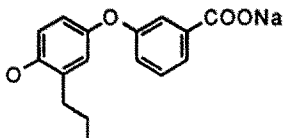
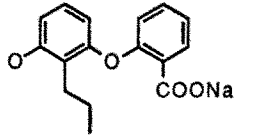
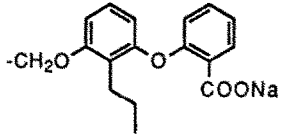
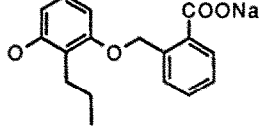
Scheme II.

a) (i) NaH/DMF, (ii) methyl 2-(bromomethyl)benzoate (21%).

Scheme III.

a) 1-bromo-3-chloropropane/ K_2CO_3 /DMF (92%); b) 2,4-dihydroxy-5-ethylacetophenone/ K_2CO_3 /KI/2-butanone/DMSO (54%); c) aq. NaOH/THF/MeOH (70%).

Table I. Inhibition of Specific Binding of ^3H -LTB₄

			
Compound	R	Human Neutrophils (IC ₅₀ , nM) ^a	Guinea-pig Lung Membranes (K _i , nM) ^a
LY255283	—	85 ± 7.9 ³	77 ± 9.4
SC-41930	—	200 ^{4b}	15 ± 3.0
11		460	140 ± 36
12		190	90 ± 20
13		290	130 ± 30
14		330	1200 ± 370
15		160	256 ± 38
16		24	39 ± 11
17		94	59 ± 12
18		130	244 ± 113
LTB ₄	—	1.9 ± 0.050 ¹⁰	0.12 ± 0.015

function for compound 18, was synthesized through monoalkylation of 2-propylresorcinol (2) as shown in Scheme II. Final coupling of the diaryl ether phenols or phenol 8 to the hydroxyacetophenone moiety is exemplified in Scheme III for the synthesis of compound 16, and was effected with standard methodology. All compounds were converted to their sodium salts and purified via reversed-phase liquid chromatography.⁷

Initial compounds of the series (11-13) proved disappointing in terms of binding affinity for both human neutrophil⁸ and guinea-pig lung membrane⁹ (Table I). Clearly the presence of a lipophilic group within close proximity to the acid binding function of both LY255283 and SC-41930 is critical for maximum activity, and the gem dimethyl unit of LY255283 could well be playing an analogous role to that of the *n*-propyl group of SC-41930.⁵ While little change was observed in binding affinity with *n*-propyl-substituted derivatives 14 and 15, compound 16, the *n*-propyl derivative of meta/ortho diaryl ether 13, resulted in a significant gain in activity. This exercise underscores the sensitive nature of the LTB₄ receptor acid-binding pocket to adjacent lipophilicity and small geometrical permutations in the acid portion of a given series of antagonists. Two attempts were made to refine the binding of 16 through the insertion of a methylene spacer, first in the chain connecting the diaryl ether to the hydroxyacetophenone moiety (compound 17), and then between the diaryl ether oxygen and the acid-bearing phenyl group (compound 18). Both modifications resulted in a loss in binding affinity.

Compounds 16, 17, and 18 were further evaluated as to their ability to inhibit LTB₄-induced chemotaxis of human neutrophils,¹¹ an important assay when considering the potential antiinflammatory role of LTB₄ antagonists. Compound 16 exhibited the greatest activity with an IC₅₀ of $0.79 \pm 0.13 \mu\text{M}$ (Figure 1), a value which compares favorably to the inhibitory range reported for SC-41930 ($1\text{--}5 \mu\text{M}$).^{4b} As predicted from inspection of neutrophil binding activities, compound 17 was slightly less active than 16, with an IC₅₀ of $3.1 \pm 0.70 \mu\text{M}$ (Figure 2). Surprisingly, compound 18 also fell in the range of SC-41930, with an IC₅₀ of $1.2 \pm 0.34 \mu\text{M}$ (Figure 3).

Figure 1. Inhibition of LTB₄-induced Chemotaxis of Human Neutrophils by Compound 16.

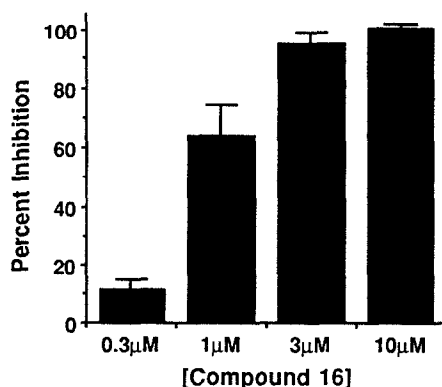


Figure 2. Inhibition of LTB₄-induced Chemotaxis of Human Neutrophils by Compound 17.

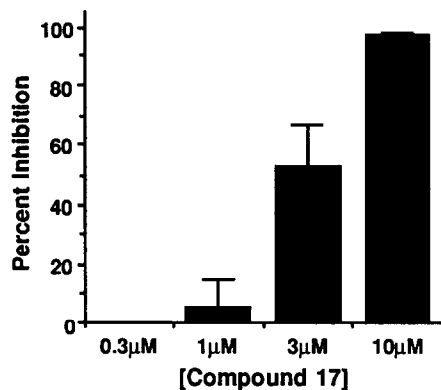
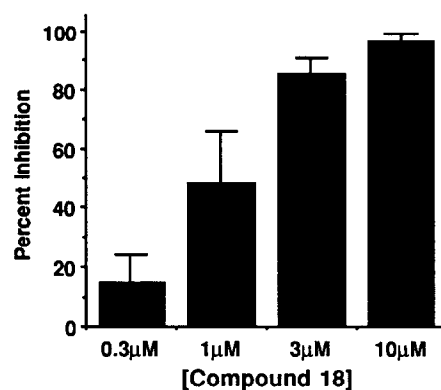


Figure 3. Inhibition of LTB₄-induced Chemotaxis of Human Neutrophils by Compound 18.



In conclusion, we have developed a short series of hydroxyacetophenone/diaryl ether LTB₄ receptor antagonists. Compound 16 (LY285009)¹² appears to take advantage of some of the major features of LY255283 and SC-41930, including an out-of-plane disposition of an acidic functionality in close proximity to a secondary lipophilic binding group. This new achiral antagonist has a binding affinity for human neutrophils approximately 8-fold higher than the reported value for SC-41930, and is approximately 3-fold more potent than LY255283. We have also demonstrated potent inhibition of LTB₄-induced chemotaxis of human neutrophils with 16. Further exploration of the SAR of this series is in progress.

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- (7) Satisfactory spectral and analytical data were obtained for all final compounds.
- (8) Assay conditions are described in reference 3. For each compound, an inhibition response study was done in triplicate on cells from a single individual. Standard errors for the IC₅₀ values of these compounds can be estimated from standard deviations of IC₅₀ values for six reference LTB₄ receptor antagonists whose effects were measured on cells from five individuals. The average standard deviation was 15 ± 4% of the mean IC₅₀.
- (9) For assay conditions see: Silbaugh, S. A.; Stengel, P. W.; Cockerham, S. L.; Roman, C. R.; Saussy, D. L., Jr.; Spaethe, S. M.; Goodson, T.; Herron, D. K.; Fleisch, J. H. *Eur. J. Pharmacol.* **1992**, *223*, 57.
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- (11) For assay conditions see reference 2j.
- (12) Characterization of **16**: ¹H-NMR (DMSO-d₆) 7.57 (s, 1H), 7.41 (dd, J = 7, 2 Hz, 1H) 7.08 (t, J = 8 Hz, 1H), 6.96 (m, 2H), 6.64 (d, J = 8 Hz, 1H), 6.57 (d, J = 8 Hz, 1H), 6.52 (s, 1H), 6.26 (d, J = 8 Hz, 1H), 4.19 (t, J = 6 Hz, 2H), 4.12 (t, J = 6 Hz, 2H), 2.58 (t, J = 7 Hz, 2H), 2.52 (s, 3H), 2.50 (m, 2H), 2.19 (quintet, J = 6 Hz, 2H), 1.42 (hexet, J = 7 Hz, 2H), 1.07 (t, J = 8 Hz, 3H), 0.79 (t, J = 7 Hz, 3H); MS-FAB (m/e) 538 (p + 2, 28), 537 (p + 1, 82), 545 (100).