the C-2 hydroxyl oxygen (O(3)) of topa in complex 1 is slightly longer than the above value, but Cu(II) can promote the oxidation of benzylamine. Although these preliminary experiments are not sufficiently extensive to be able to define the mechanism, the finding that Cu(II) is particularly essential for the acceleration of the oxidation of benzylamine is of interest in connection with the mechanism of amine oxidases. Complex 1 might provide significant information on the catalytic roles of the topa residue and copper in copper-containing amine oxidases. Further studies on the spectral and redox properties and the detailed catalytic reaction of complex 1 are in progress.

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Supplementary Material Available: Figure showing the stereoscopic view of the molecular packing for the complex 1 and tables of positional parameters and isotropic temperature factors for non-hydrogen atoms, interatomic distances and angles, positional parameters and isotropic temperature factors for hydrogen atoms, and anisotropic thermal parameters for non-hydrogen atoms (6 pages); listings of observed and calculated structure factors (9 pages). Ordering information is given on any current masthead page.

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Novel Tight-Binding Inhibitors of Leukotriene A₄ Hydrolase

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Leukotriene (LT) A_4 hydrolase (EC 3.3.2.6)¹ is a zinc-containing² monomeric enzyme (MW \approx 70 kDa) which catalyzes the formation of LTB₄ (5(S),12(R)-dihydroxy-6,14-cis-8,10-transeicosatetraenoic acid) from its natural substrate LTA_4 ((5S)-5,6-oxido-7,9-trans-11,14-cis-eicosatetraenoic acid),^{3,4} one of the Scheme I. Proposed Mechanism for the Hydrolysis of LTA. Catalyzed by LTA₄ Hydrolase



Scheme II. Proposed Mechanism for the Aminopeptidase Activity of LTA₄ Hydrolase







^aAll assays were performed in Tris-HCl buffer (0.05 M, pH 8.0) with L-alanyl p-nitroanilide (1.5 mM) as substrate unless otherwise indicated. LTA_4 hydrolase (1.4 µg) purified from human leukocytes was added for each assay (final volume = 1.0 mL). p-Nitroaniline formation was monitored spectrophotometrically at 405 nm. Dixon plot was used to determine the K_i value. ^bLess than 50% inhibition was observed at that concentration. 'Slow binding behavior was observed. The inhibitor-enzyme mixture was incubated for 30 min prior to addition of the substrate.

physiologically important processes in the arachidonic acid biosynthetic pathway.¹ It also catalyzes the hydrolysis of some L-amino acid amides.^{3,4} The mechanisms of both enzymatic reactions, though not being elucidated, are believed to involve a general base (from a carboxylate) and a Lewis acid (from Zn^{2+}) (Schemes I and II). The zinc ion may coordinate to the nucleophilic water molecule to facilitate the general base catalysis. The peptidase and epoxide hydrolase activities, though they occur in the same active site,³ may use a different general base (a carboxylate residue) as indicated in recent site-directed mutagenesis studies.5

Since LTB₄ is a proinflammatory mediator which stimulates adhesion of circulating neutrophils to vascular endothelium and directs their migration toward sites of inflammation, it is of interest to develop inhibitors of LTA₄ hydrolase as potential antiinflam-

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amides such as L-Ala *p*-nitroanilide ($K_m = 0.5 \text{ mM}$, $V_{max} = 530 \text{ nmol/min/mg}$ enzyme) and L-Pro *p*-nitroanilide ($K_m = 0.1 \text{ mM}$, $V_{max} = 130 \text{ nmol/min/mg}$ enzyme) are good substrates. The Denantiomers are not substrates. Values of K_m and V_{max} in the ranges 7-30 μ M and 1.7-3.0 μ mol/mg/min, respectively. tively, have been reported using LTA₄ as substrate: Radmark, O.; Haeggs-trom, J. Adv. Prostaglandin, Thromboxane, Leukotriene Res. 1990, 20, 35-45. The complementarity components were designed on the basis of computer modeling using MACROMODEL.

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Scheme III. Proposed Inhibitor-Enzyme Complex



matory agents. LTA_4 and its analogs (e.g., LTA_3 and LTA_5)⁶ are irreversible inhibitors, and some inhibitors of other Zn²⁺containing amino peptidases and angiotensin-converting enzymes are also reversible inhibitors of LTA₄ hydrolase,⁷ supporting the proposed mechanism.

Our previous studies on the inhibition of this enzyme with more than 10 peptide-based, synthetic transition-state analog inhibitors (including α -hydroxy β -amino acids and their peptide derivatives, fluoro ketone and phosphoramidate derivatives)⁴ have led us to develop another class of compounds (see Table I, 6-8) which have proven to be better inhibitors than those simply based on the amidase activity. These inhibitors contain a transition-state mimic of the enzyme-catalyzed amide cleavage as a "core" and additional complementarity components (the aromatic moieties) which resemble the hydrophobic nature of the conjugated polyene system of the natural substrate LTA₄, which binds to the enzyme more tightly than the amide substrates.⁴ We chose α -keto esters instead of α -keto amides⁸ for further development because the ester derivative 2 binds to the enzyme more tightly than the amide 1. The α -keto amide with a free carboxyl group (4) is, however, a better inhibitor (IC₅₀ = 0.5 μ M) than the corresponding α -(S)-OH derivative (IC₅₀ = 20 μ M; the α -(R)-OH derivative is not an inhibitor).4

Further adjustment of the inhibitor structure at the P1' and P2-P3 sites led to the development of an α -keto β -amino ester (8) with $K_i = 0.1 \ \mu$ M. The NMR spectra indicate that both α -keto β -amino amide and α -keto β -amino esters are completely hydrated in water and 60% hydrated in DMSO containing 5% H_2O . We therefore propose that the inhibitor exists as a gem-diol bound in the enzyme active site;9 the free amino group and one of the hydroxyl groups may coordinate to the Zn²⁺ (as N-Boc and N-Cbz derivatives are not inhibitors) and the other hydroxyl group interacts with the general base (CO_2^-) via H-bonding (Scheme III).

In summary, the α -keto β -amino esters developed in this study are a new class of selective¹⁰ inhibitors of LTA₄ hydrolase. The bound inhibitor seems to resemble the transition-state structure of the enzymatic amide cleavage and LTA₄ binding.¹² Work is in progress to determine the structure of the inhibitor-enzyme complex for mechanistic investigation and to develop better inhibitors.

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Supplementary Material Available: Listings of experimental procedures and physical data (mass and NMR data) for all of the inhibitors (5 pages). Ordering information is given on any current masthead page.

(12) The procedures for the synthesis of compounds in Table I are essentially the same as described previously.⁴ The keto esters or keto amides were prepared from the corresponding OH compounds via Swern oxidation.

Substituent Effects on Diazomethanes and Diazirines by ab Initio Molecular Orbital Calculations

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Diazo compounds $(1)^1$ and the isomeric diazirines $(2)^2$ are important substrates in synthetic and mechanistic studies, particularly as precursors of carbenes,³ and substituent effects on all these species is a topic of major current interest.¹⁻³ Although many substituted derivatives of these compounds have been prepared, there is considerable uncertainty^{1,2} as to how substituents affect the ground-state stability of 1 and 2.4 The effect of substituents on the diazomethane/diazirine equilibrium^{4b-e} and on reactions of these species such as carbene formation cannot be properly evaluated in the absence of an understanding of ground-state substituent effects on 1 and 2.

We have recently reported ab initio molecular orbital studies on the effect of substituents on ketenes (3),^{5a} which are isoelectronic to diazomethanes, and on diazocyclopolyenes such as diazocyclopentadiene (4),⁵⁶ for which there is evidence for aromatic

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