2-(Peptidamido)-cyclobutanones: A Novel Strategy for the Inhibition of Serine Elastases

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The synthesis of 2-peptidamidocyclobutanones is described as a novel class of competitive elastase inhibitors. Their inhibitory potency, although modest (i.e., micromolar range), is up to two orders of magnitude better than their acyclic analogues. The enhanced potency is explained, in part, by the tendency of an sp^2 hybridized cyclobutyl carbonyl to release internal ring strain by conversion to an sp^3 hybridized center. © 1992 Academic Press, Inc.

INTRODUCTION

Elastolytic enzymes of the serine proteinase family have received considerable attention recently (1, 2), especially as targets for inhibitor design (3). This effort has been based upon a general acceptance of the proteinase-antiproteinase hypothesis (4) and a good understanding of the mechanism of serine proteinase catalysis (5-7).

The so called "transition state analogues" of human neutrophil elastase (HNE) rank as the most potent competitive inhibitors of this enzyme yet discovered (8). Peptidyl trifluoromethyl ketones (9) and aldehydes (10) inhibit HNE in the nanomolar range. For many transition state analogues a carbonyl functionality replaces the scissile amide linkage of a good substrate (9). Initial binding is followed by a covalent interaction between the carbonyl of the inhibitor and the hydroxyl of the active site serine. Peptidyl aldehydes (11) and peptidyl trifluoromethyl ketones (12) form hemiacetal and hemiketal-like structures, respectively, upon interaction with serine proteinases. These tetrahedral species are tightly bound because they mimic a high energy intermediate of the hydrolytic pathway.

Peptidamidoalkyl ketones, in contrast to aldehydes, are sterically encumbered at the carbonyl and are generally poor inhibitors of elastase (13). Also, they lack the powerful electron withdrawing groups of the trifluoromethyl ketones and related inhibitors such as esters (14) and diketones (15). In this work we introduce an alternate strategy, the incorporation of internal ring strain (16), for the inhibition of serine proteinases. This is the first instance in which the inherent strain of a cyclobutanone has been exploited for the design of a proteinase inhibitor. A

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related effort on bicyclo[3.2.0]heptan-6-one synthesis led to the discovery of time-dependent β -lactamase inhibitors (17).

EXPERIMENTAL

Computer Modeling

All calculations were done on a Silicon Graphics 4D25GT personal IRIS workstation running the Polygen Quanta Rev 2.1A graphics package with CHARMm Rev. 21 molecular minimization/dynamics code and AMPAC Rev 1.00.

Coordinates for the model compounds were calculated from a 2D model with the Chemnote molecule sketching/building package. Partial atomic charges were from an INDOCL calculation. The global energy minimum was found with a grid search in torsional angle space. A grid of 5° was applied around the appropriate torsional angles and an adopted basis Newton Raphson minimization based on the CHARMm parameter set. The coordinates of the lowest energy structure were then used as input into an AMPAC calculation. This calculation employed full geometry optimization and a 100× precision factor on the convergence criterion. Parallel calculations were done for each of the AM1, MNDO, and MINDO3 Hamiltonians.

Synthesis

2-(N-Benzyloxycarbonyl-L-alanyl-L-alanyl-L-prolinamido)-cyclobutanol N-Benzyloxycarbonyl-L-proline (0.33 g, 1.32 mmol) was dissolved in dry tetrahydrofuran (25 ml), blanketed with argon, and cooled in a dry ice/acetone bath. The solution was treated dropwise with isobutyl chloroformate (154 µl, 1.19 mmol) and N-methylmorpholine (128 μ l, 1.16 mmol) and stirred for 10 min. The resulting mixture was treated with a solution of trans-2-aminocyclobutanol in dry tetrahydrofuran (10 ml) and stirred at room temperature for 4 h. Ethyl acetate (200 ml) was added and the organic layer was washed with 5% HCl (1 \times 50 ml), saturated sodium bicarbonate solution (1 \times 50 ml), and saturated sodium chloride solution $(1 \times 50 \text{ ml})$ followed by drying over sodium sulfate. The solvent was evaporated and the residue was applied to a silical gel column. The 2-(N-benzyloxycarbonyl-L-prolinamido)cyclobutanol was eluted with MeOH/CH₂Cl₂ (1:9) under pressure. Evaporation gave **1** as a syrup (0.165 g, 44.7%) $(R_f 0.35; \text{ ir } 3305, 2972, 1687, 1656, 1656)$ 1421, 1357, 1120, 735 cm⁻¹; ¹H NMR d 1.60–2.35 (m. 8H, H-2, H-2', H-3, H-3', NCH₂(CH₂)₂CH), 3.4–4.0 (m, 5H, H-1, H-2, NCH₂(CH₂)₂CH, OH), 4.31 (bs, 1H, NCH₂(CH₂)₂CH), 5.16 (m, 2H, CH₂Ph), 6.24, 7.16 (two d, 1H, NH), 7.36 (s, 5H, CH₂Ph).

Syrupy 1 was dissolved in methanol (25 ml) blanketed with argon and treated with 10% palladium on carbon. The argon gas was exchanged for hydrogen (3 atm) and the reaction mixture was stirred at room temperature for 4 h. The contents of the reaction flask were filtered, the cake was washed with MeOH (2×30 ml), and the filtrate was concentrated to a syrupy residue. A portion of the residue (0.045 g, 0.19 mmol) was dissolved in dry tetrahydrofuran (7 ml) to make a solution of 2.

N-Benzyloxy-carbonyl-L-alanyl-L-alanine (0.048 g, 0.20 mmol) was dissolved in dry tetrahydrofuran, blanketed with argon, and cooled in a dry ice/acetone bath. The resulting solution was treated dropwise with isobutylchloroformate (26 μ l, 0.20 mmol) and N-methylmorpholine (22.6 μ l, 0.21 mmol) and stirred for 10 min. The reaction mixture was treated with 2, in solution, and warmed to room temperature for 4 h.

Ethyl acetate (100 ml) was added and the organic layer washed with 5% HCl (1 × 25 ml) and saturated sodium bicarbonate solution (1 × 25 ml) and dried over sodium sulfate. The organic layer was filtered, evaporated to dryness, and chromatographed under pressure on silica gel using MeOH/CH₂Cl₂ (1:9). 2-(N-Benzyloxycarbonyl-L-alanyl-L-prolinamido)cyclobutanol, **3**, was isolated as a syrup (0.036 g, 28%): R_f 0.23; ir 3297, 1703, 1656, 1639, 1544, 1460, 1250, 735 cm⁻¹; ¹H NMR d 1.25–1.4 (m, 6H, CHCH₃, CHCH₃), 1.5–2.6 (m, 6H, H-3, H-3', H-4, H-4', NCH₂(CH₂)₂CH), 3.45–3.95, 4.0–4.2 (two m, 5H, H-1, H-2, NCH₂(CH₂)O₂CH, OH), 4.25–4.6, 4.83 (m, bs, 3H, CH₃CH, CH₃CH, NCH₂(CH₂)₂CH), 5.10 (s, 2H, CH₂Ph), 5.43, 5.69 (two bs, 1H; NH), 7.2–7.5 (bs, 6H, CH₂Ph), 7.89, 7.99 (two bs, 1H, NH); HRMS m + 1/e calcd. for C₂₃H₃₂N₄O₆: 461.2400. Found: 461.2406.

2-(N-Benzyloxycarbonyl-L-alanyl-L-prolinamido)-cyclobutanone (4a, 4b). Dimethyl sulfoxide (208 μ l, 2.93 mmol) was added dropwise to a solution of oxalyl chloride (128 μ l, 1.46 mmol) at about -78° C in dry dichloromethane (7 ml). The resulting solution was stirred for 30 min then treated dropwise with a solution of 2-(N-benzyloxycarbonyl-L-alanyl-L-alanyl-L-prolinamido)cyclobutanol, 3, (0.021 g, 0.045 mmol) in dry CH₂Cl₂ (7 ml). The reaction mixture was stirred an additional 30 min at -78° C, then treated with triethylamine (407 μ l, 2.9 mmol) and warmed to room temperature for 1 h. The reaction mixture was diluted with ethyl acetate (40 ml) then washed with 1 n HCl (10 ml), saturated bicarbonate solution (2 × 10 ml) and brine (3 × 20 ml). The organic layer was dried (Na₂SO₄) and concentrated, the residue was subjected to flash chromatography (MeOH/CH₂Cl₂, 1:9) to yield the title compound 4 as a colorless syrup (0.019 g, 92%, R_f 0.35). Separation by HPLC (EtOAc/hexane, 3:1, 4 ml/min, Waters 7.6 mm × 30 cm μ Porasil, 10 μ m) afforded two fractions.

Short retention isomer 4a: $R_1 = 23.5$ min; ir 3299, 1793, 1638, 1532, 1460, 1244, 736, 699 cm⁻¹; ¹H NMR d 1.29, 1.35 (two d, 6 J = 6.5, 6.8, $2 \times \text{CH}_3\text{CH}$), 1.85–2.30 (m, 6, NCH(CH₂)CH, H-3, H-3'), 2.83 (m, 2, H-4, H-4'), 3.6–3.8 (m, 2, NCH₂ (CH₂)₂CH, 5.10 (s, 2, PhCH₂), 5.64 (d, 1, J = 7, NH), 7.35 (s, 5, PhCH₂), 7.60, 7.86 (two d, 2, J = 7.2, 7.1, $2 \times \text{NH}$); HRMS m + 1/e calcd. for $C_{23}H_{30}N_4O_6$: 459.2244. Found: 459.2245.

Long retention isomer **4b**: $R_t = 26$ min; ir 3303, 1791, 1639, 1532, 1460, 1244, 736, 700 cm⁻¹; ¹H NMR d 1.28, 1.35 (two d, 6, J = 6.6, 6.6, $2 \times \text{CH}_3\text{CH}$), 1.7–2.2 (m, 6, NCH(CH₂)CH, H-3, H-3'), 2.84 (m, 2, H-4, H-4'), 3.6–3.8 (m, 2, NCH₂ (CH₂)₂CH), 4.5 (m, 2, $2 \times \text{CH}_3\text{CH}$), 4.8–4.9 (m, 2, NCH₂(CH₂)₂CH, H-2), 5.09 (s, 2, PhCH₂), 5.65 (d, 1, J = 6.9, NH), 7.35 (s, 5, PhCH₂), 7.67 (d, 1, J = 7, NH), 7.92 (bs, 1, NH); HRMS m + 1/e calcd. for C₂₃H₃₀N₄O₆: 459.2244. Found: 459.2240. 3-(N-Benzyloxycarbonyl-L-alanyl-L-alanyl-L-prolinamido)-2-butanone (8a, 8b). N-Benzyloxycarbonyl-L-proline (0.249 g, 1.0 mmol) was treated, as described for

1, with isobutylchloroformate (137 μ l, 1.06 mmol), *N*-methylmorpholine (99 μ l, 1.06 mmol) and threo-3-amino-2-butanol (0.090 g, 1.0 mmol). Standard workup and flash chromatography on silica gel (MeOH/CH₂Cl₂, 1:9) yielded the amidoalcohol **5** (0.265 g, 82.8%, R_f = 0.34); ir 3331, 2975, 1701, 1686, 1420, 1357, 1121, 736 cm⁻¹; ¹H NMR d 0.95–1.25 (m, 6, CH₃CHCHCH₃), 1.85–2.60 (m, 5, CH₂CH₂, OH), 3.55 (bs, 2, NCH₂), 3.65, 3.85 (2bs, 2, CH₃CHCHCH₃), 4.33 (m, 1, NCHCO), 5.16 (s, 2, CH₂Ph), 6.10, 6.60 (2 bs, 1, NH), 7.35 (s, 5, CH₂Ph).

Amidoalcohol 5 (0.260 g, 0.81 mmol) was dissolved in ethanol (125 ml) treated with 10% Pd/C (25 mg) and worked up as described for 2 to give syrupy 6 (0.150 g, 97.5%), which was used without further purification.

N-Benzyloxycarbonyl-L-alanyl-L-alanine (0.119 g, 0.404 mmol) was dissolved in tetrahydrofuran (10 ml) treated with isobutylchloroformate (53 μl, 0.409 mmol), *N*-methylmorpholine (46 μl, 0.418 mmol) and **6** (0.075 g, 0.403 mmol) as previously described. Workup and flash chromatography (MeOH/CH₂Cl₂ 1:19 to 1:9) gave 2-(*N*-benzyloxycarbonyl-L-alanyl-L-prolinamido)cyclobutanol **7** (0.10 g, 53.7%, $R_f = 0.25$); ir 3305, 2975, 1657, 1638, 1536, 1455, 1265, 735 cm⁻¹; ¹H NMR d 1.04–1.12 (m, 6, CH₃CHCHCHCH₃), 1.25, 1.33 (2 d, 6, CH₃CH, CH₃CH), 1.9–2.40 (m, 4, NCH₂(CH₂)₂CH), 3.81 (d, 1, OH), 3.5–3.9 (m, 4, NCH₂(CH₂)₂CH, CH₃CHCHCHCH₃), 4.3–4.65, 4.84 (m, bs, 3, CH₃CH, CH₃CH, NCH₂(CH₂)₂CH), 5.94 (ABq, 2, CH₂Ph), 5.80 (bs, 1, NH), 7.00, 7.09 (2 d, 1H, NH), 7.33 (s, 5, CH₂Ph), 8.05 (d, 1, NH).

Amido butanol 7 (0.09 g, 0.195 mmol) was treated as described for 3 with dimethyl sulfoxide (800 μ l, 11.3 mmol), oxalyl chloride (493 μ l, 5.65 mmol), and triethylamine (1.60 ml, 11.5 mmol) to give, after chromatography (MeOH/CH₂Cl₂, 1:19 to 1:9), the title compound 8 (0.060 g, 66.4%, $R_f = 0.33$) as a mixture of epimers at C-3. This material was separated into two fractions by HPLC as described above.

Short retention isomer **8a**: $R_t = 23$ min; ir 3303, 2926, 1721, 1639, 1530, 1459, 1250, 737 cm⁻¹; ¹H NMR d 1.26–1.38 (m, 9, CH₃CHCHOCH₃, CHCH₃, CHCH₃), 1.9–2.3 (m, 4, NHCH₂(CH₂)₂CH), 2.13 (s, 3, CH₃CHCOCH₃), 3.55–3.80 (m, 2, NCH₂(CH₂)₂CH), 4.35–4.65 (m, 3, CH₃CH, CH₃CH, CH₃CHCOCH₃), 4.83 (m, 1, NCH₂(CH₂)₂CH), 5.11 (m, 2, CH₂Ph), 5.62 (bs, 1H, NH), 7.34 (s, 5, CH₂Ph), 7.44 (m, 1, NH), 7.63 (bs, 1, NH); HRMS m + 1/e calcd. for C₂₃H₃₃N₄O₆: 461.2400. Found: 461.2402.

Long retention isomer **8b**: $R_t = 30$ min; ir 3283, 2927, 1720, 1640, 1459, 1248, 1028, 700 cm⁻¹, ¹H NMR d 1.24–1.38 (m, 9, CH₃CHCOCH₃, CH₃CH, CH₃CH), 2.0–2.25 (m, 4, NHCH₂(CH₂)₂CH), 2.18 (s, 3, CH₃CHCOCH₃), 3.6–3.8 (m, 2, NCH₂(CH₂)₂CH), 4.4–4.55 (m, 3, CH₃CH, CH₃CH, CH₃CHCOCH₃), 4.82 (m, 1, NCH₂(CH₂)₂CH), 5.12 (m, 2, CH₂Ph), 5.58 (d, 1, NH), 7.35 (bs, 6, CH₂Ph, NH), 7.65 (d, 1, NH); HRMS m + 1/e calcd. for C₂₃H₃₂N₄O₆: 461.2400. Found: 461.2402.

3-(N-Benzyloxycarbonyl-L-alanyl-L-prolinamido)-1,1,1-trifluoro-2-butanone (12). A solution of N-benzyloxycarbonyl-L-proline (0.031 g, 0.13 mmol) in anhydrous tetrahydrofuran was treated with isobutyl chloroformate (16.3 μ l, 0.3 mmol), N-methylmorpholine (13.8 μ l, 0.13 mmol) and a solution of 3-amino-1,1,1-trifluorobutanol (0.020 g, 0.14 mmol) in tetrahydrofuran (5 ml) as described above.

Standard workup yielded a residue: ir 3295, 1681, 1666, 1174, 1138, 734, 696 cm⁻¹. The residue was purified by flash chromatography (MeOH/CH₂Cl₂, 1:19) to give **9** (0.023 g, 48%) in two fractions.

High R_f fraction: 0.65 (HOAc/MeOH/CH₂Cl₂, 0.1:1:9); ¹H NMR d 1.25 (s, 3, CH₃), 1.9–2.3 (m, 4, NHCH₂(CH₂)₂CH), 3.53 (bs, 2, NCH₂(CH₂)₂CH), 4.1 (s, 1, OH), 4.22 (s, 2, H-2, H-3), 4.67 (s, 1, NCH₂(CH₂)₂CH), 5.14 (s, 2, PhCH₂), 6.67 (bs, 1, NH), 7.35 (s, 5, PhCH₂).

Low R_f fraction: 0.61 (HOAc/MeOH/CH₂Cl₂, 0.1:1:9); ¹H NMR d 1.26 (s, 3, CH3), 1.9–2.3 (m, 4, NHCH₂(CH₂)₂CH), 3.52 (s, 2, NCH₂(CH₂)₂CH), 3.85 (s, 1, OH), 4.0–4.4 (m, 2, H-2, H-3), 4.8 (s, 1, NCH₂(CH₂)₂CH), 5.14 (s, 2, PhCH₂), 6.15–7.1 (bm, 1, NH), 7.35 (s, 5, PhCH₂).

A solution of **9** (combined fractions, 0.057 g, 0.15 mmol) in absolute MeOH (10 ml) was treated with 10% Pd/C (20 mg) and hydrogen gas as previously described to give **10** as a residue ir 3325, 1625, 1525, 1175, 1142 cm⁻¹.

A partial suspension of benzyloxycarbonyl-L-alanyl-L-alanine (0.040 g, 0.136 mmol) in dry tetrahydrofuran was treated with isobutylchloroformate (17.6 μ l, 0.136 mmol) and *N*-methylmorpholine (14.9 μ l, 0.136 mmol) and 3-(L-prolinamido)-1,1,1-trifluorobutan-2-ol, **10** (0.36 g, 0.15 mmol). After stirring at room temperature 3 h, 1-ethyl-3-(3-dimethylaminopropyl)-carbodiimide · HCl (0.013 g, 0.068 mmol) was added and the suspension was stirred overnight at room temperature. Standard workup gave a residue which was purified by flash chromatography (MeOH/CH₂Cl₂, 1:19–1:9) to yield **11** as a solid (0.052 g, 74%): R_f 0.48, 0.44 (MeOH/CH₂Cl₂, 1:9): ir 3305, 1704, 1635, 1270, 1176, 1143, 737, 699 cm⁻¹; ¹H NMR d 1.1–1.4 (m, 9, CH₃), 2.0–2.1 (m, 4, NCH₂(CH₂)₂CH), 3.61, 3.74 (m, 1, NCH₂(CH₂)₂CH), 3.9–4.5 (m, 3, 2 × CHCH₃, H-2), 4.82 (d, 1, J = 7.0, NCH₂ (CH₂)₂CH), 5.10 (s, 2, PhCH₂), 5.45 (bs 1, H-2), 5.74 (d, 1, J = 7.7, NH), 7.34 (s, 5, PhCH₂), 7.56 (s, 1, NH), 8.09 (d, 1, J = 7.6, NH).

The amido trifluoromethylbutanol **11** was treated as described above with oxalyl chloride (140 μ l, 1.60 mmol), dimethyl sulfoxide (230 μ l, 3.24 mmol), and triethylamine (450 μ l, 3.23 mmol). Workup followed by flash chromatography (MeOH/CH₃Cl₂, 1:9) gave the title **12** as a solid (0.024 g, 72%): R_f 0.60–0.67; ir 3302, 1697, 1634, 1266, 1182, 738, 702 cm⁻¹; ¹H NMR d 1.31 (m, 9, CH₃), 1.9–2.2 (m, 3, 2 × CH₃CH, H-2), 3.6–3.8 (m, 2, NCH₂(CH)₂CH), 4.3–4.6 (m, 3, 2 × CH₃CH H-2), 4.82 (m, 1, NCH₂(CH₂)₂CH), 5.10 (Abq, J = 12, 1, PhCH₂) 5.55 (d, 1, J = 7.9, NH), 7.35 (s, 5, PhCH₂), 7.6 (m, 2, NH); HRMS m + 1/e calcd: for C₂₃H₂₉N₄O₆F₃: 515.2118. Found: 515.2115. A sample of this material was further purified by HPLC; one diffuse peak with $R_I = 14.8$ minutes was recovered.

Elastase Inhibition Assay

The compounds were assayed for inhibitory activity against human neutrophil elastase obtained from Elastin Products (lot SE 563) and porcine pancreatic elastase from Boehringer-Manheim, West Germany (lot 11111223-02). The elastase inhibition assay employed the following: buffer, 1.0 m NaCl, 0.2 m Hepes, pH 7.5; substrate, methoxysuccinyl-L-alanyl-L-alanyl-L-prolyl-L-valyl-4-nitroanalide dissolved in DMSO; test compound, dissolved in DMSO and elastase dissolved in

Inhibitor	Isomer	K_i (mм)	
		HNE	PPE
3 Z—A—A—P—NHOH	_	2.0	0.85
4a Z—A—A—P—NH	Short retention	0.55	0.025
4b Z—A—A—P—NH	Long retention	1.2	ND
8a Z—A—A—P—NH	Short retention	>2.5	>2.0
8b Z—A—A—P—NH O O 12 Z—A—A—P—NH CF ₃	Long retention	>5.0	1.6
12 Z—A—A—P—NH CF ₃	_	0.005	0.0001

TABLE 1
Inhibition Constants of Elastase Inhibitors

buffer. Assays were monitored at 410 nm in a 25°C, temperature-stabilized six-position Beckman DU-40 spectrophotometer.

To conduct the assay, the following were mixed in a 500- μ l quartz cuvette: 20 μ l buffer; 150 μ l distilled, deionized water; 20 μ l substrate; 20 μ l inhibitor; and 10 μ l elastase. Absorbance was monitored for 30 min and the slope of the linear range was calculated. For the purpose of calculating an inhibition constant, these steps were repeated for the following solution conditions: elastase, 5–20 nm; substrate, 0-500 μ M; and inhibitor, 0–2000 μ M. The inhibition constant, K_i , was then calculated from the Dixon plot (1/velocity vs [I]) for each concentration of substrate. The K_i s obtained are presented in Table 1.

RESULTS AND DISCUSSION

Nucleophilic addition to aldehydes and ketones has been chosen as a model system for understanding the inhibitory potential of peptidamido ketones and aldehydes (18). For example, trifluoromethyl ketones are extensively hydrated in aqueous systems (i.e., low $K_{\rm H_2O}$), alkyl ketones are not (19). This correlates with the inhibitory potency of their peptidyl analogues, even though a linear relationship between K_i and $K_{\rm H_2O}$ cannot be demonstrated (18).

Cyclobutanones are generally more susceptible to nucleophilic attack than their

Compound	Hamiltonian	Heat of formation (H_f) (kcal/mol)	$\Delta H_f^{\ u}$ (kcal/mol)	$\Delta \Delta H_f^b$ (kcal/mol)
NH ₂ AM1 MNDO MINDO3		-50.9 -465		
	-463 -53.5			
NH ₂ AMI MNDO MINDO3	-122.3	71.4		
	MNDO	-104.8	58.3	
	MINDO3	-108.5	55.0	
NH ₂ O AM1 MNDO MINDO3	AM1	-21.7		
	MNDO	-31.1		
	-39.1			
MI	AM1	-97.2	75.5	4.1
	MNDO	-97.5	66.4	8.1
	MINDO3	-102.1	63.0	8.0

TABLE 2

AMPAC Calculations on Model Compounds

acyclic counterparts. A tentative result (20) indicates that the four-membered ring is significantly more hydrated than its acyclic counterpart. Moreover, the equilibrium value for thiol addition to cyclobutanone, K(RSD) is about 20 times the value for acetone. Reaction rates are also affected; the reduction of cyclobutanone by sodium borohydride is about 500 times faster than it is for open chain ketones (21).

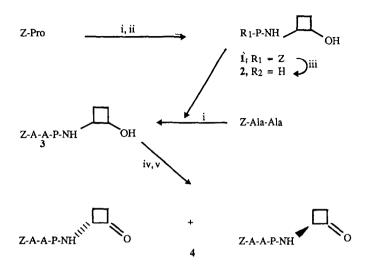
These phenomena are attributed to the strain introduced by incorporating an sp^2 -hybridized carbonyl (120° preferred) into a four-membered ring (90° angles). Some of this "I strain" (16) is released by conversion to an sp^3 hybridized center (109.5° preferred), i.e., through reduction, thiohemiketal formation, or, presumably, formation of a tetrahedral hemiketal in the enzyme active site.

The energetics of the cyclobutanone/hydrate pair were modeled and compared to their acyclic counterparts. Results of the AMPAC calculation are given in Table 2. The MNDO and MINDO3 Hamiltonians predict that the "I strain" effect would favor the hydration of the cyclobutanone over its acyclic analogue by some 8 kcal/mol. The calculation based on the AM1 Hamiltonian also favors cyclobutanone hydration but by only 4 kcal/mol. This analysis ignores any entropic effects, which should also favor the cyclobutanone hydration. The conformationally restricted four-membered ring has fewer degrees of rotational freedom than its acyclic analogue.

The above considerations led to the formulation of potential PPE inhibitors 4a, 4b in which a cyclobutanone is grafted onto peptidyl chains complementary to the elastase active site. The peptidyl backbone of 4 (A-A-P) is recognized as one of the preferred sequences for binding into the S2-S4 subsites of porcine pancreatic elastase (22). The N-blocking group, benzyloxycarbonyl (Z), was chosen for syn-

^a (H_f of ketone)-(H_f of corresponding hydrate).

^b Difference in ΔH_f between acyclic and cyclic forms.



- 4a short retention time isomer.
- 4b long retention time isomer

SCHEME 1. Synthesis of 2-peptidamidocyclobutanones: (i) NMM, isobutylchloroformate; (ii) trans-2-amino-cyclobutanol; (iii) H₂, Pd/C; (iv) DMSO, oxalylchloride, Et₃N; (v) HPLC.

thetic convenience and because it has been successfully incorporated into human neutrophil elastase inhibitors (23). Retrosynthetic analysis yielded three synthons; Z-proline and -Ala-Ala are commercially available and trans-2-aminocyclobutanol was prepared by the method of Hartmann et al. (24) (Scheme 1).

Aminocyclobutanol, 1, was coupled to Z-proline using EDCl or the mixed anhydride procedure (25); the latter gave more consistent results. The diastereomeric mixture obtained was deprotected and used for subsequent coupling to Z-Ala-Ala without further purification. This coupling reaction followed the first in all respects and gave 3 as a mixture of stereoisomers. The final oxidation step was accomplished with PCC or Swern conditions (26); the latter was cleaner and the method of choice. This protocol gave 4 as a mixture of stereoisomers at C-2, which were separated by HPLC. The stereochemistry of 4a and 4b at C-2 was not assigned.

An entirely analogous procedure, beginning with 3-amino-2-butanol (27) yielded the ketones **8a** and **8b**, which were assayed as controls against both PPE and HNE. Similarly, **12** was obtained from the previously described 3-amino-1,1,1-trifluoro-2-butanol (13) as a mixture which, in our hands, could not be separated into its constituent isomers (Scheme 2).

The compounds were tested against both PPE and HNE (see Table 1). The inhibition constants are generally much lower against PPE than HNE. PPE prefers to bind small aliphatic side chains such as alanine at the S-1 subsite; HNE

Z-pro + NH₂

$$R_2$$
 OH
 R_1 -P-NH
 R_2
 OH

5, R_1 =Z, R_2 =CH₃
 R_2
 R_3
 R_4 =Z-R₂=CF₃
 R_4
 R_5
 R_6
 R_1 =H, R_2 =CH₃
 R_6
 R_7
 R_8
 R_9 =CH₃
 R_9
 R_9

SCHEME 2. Synthesis of 2-peptidamidobutanones: (i) NMM, isobutylchloroformate; (ii) H₂, Pd/C; (iii) DMSO, oxalylchloride, Et₂N; (iv) HPLC.

12, R2=CF3

prefers larger aliphatic side chains such as valine (3, 28). The 3-methyl groups of 8a, 8b, and 12 mimic an alanine side chain. The PPE preference for cyclobutanone is less easily explained; however, the ring is small and may not be any more sterically demanding than an alanine mimic.

The striking feature, from the viewpoint of inhibitor design, is the real difference in inhibitory potency between the peptidaminocyclobutanones, $\mathbf{4a}$ and $\mathbf{4b}$ and the acyclic counterparts $\mathbf{8a}$ and $\mathbf{8b}$. Against PPE the difference in K_i between $\mathbf{4a}$ and $\mathbf{8b}$, the most potent isomers, approximates two orders of magnitude. A similar enhancement is seen against HNE, with a lower limit of an order of magnitude difference. This difference in inhibitor potency may be driven by an I strain effect which enhances hemiketal formation (sp^3 hybridization) in the four-membered ring.

However, an alternative explanation is hinted at by the weak inhibitory potency of the cyclobutanol 3. The cyclobutyl compounds can be viewed as conformationally restricted analogues of 8a and 8b. Thus entropic factors, in addition to the I-strain (enthalpy), may be contributing toward lower K_i s. This study does not allow us to deduce the relative importance of these nonexclusive rationalizations.

Even the best peptidyl cyclobutanone inhibitor is about two orders of magnitude less potent than comparable trifluoromethyl ketones. This result could have

been predicted from the hydration studies of Burkey and Fahey (19, 20) and also suggests a strategy for lowering the K_i s of the cyclobutyl series. Electron withdrawing groups at C-4 of the cyclobutyl ring would render the carbonyl more susceptible to nucleophilic attack.

We conclude that the cyclobutanone functionality can be exploited for the inhibition of serine proteinases such as elastase. The application of cyclobutanones for the inhibition of other hydrolytic enzymes is an obvious extension of this work (29). For example, tetrahedral structures have been demonstrated in the enzyme inhibitor complexes formed between peptidyl alkyl ketones and the metalloproteinase carboxypeptidase A (30). Alternatively, we are exploring the preparation of novel cysteine proteinase inhibitors based upon the ready addition of thiols to cyclobutanones (20).

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