Structure Activity Studies on Pseudo-symmetrical HIV-1 Protease Inhibitors

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Abstract: Lead compound 1, obtained from a previously reported structure-assisted design approach, was optimized to 17 using a traditional medicinal chemistry approach.

We have previously reported ¹ on the structure-assisted design, synthesis and enzyme inhibiting properties of a pseudo-symmetrical ² HIV-1 protease inhibitor 1. We now wish to report on structure-activity relationships for this series of bis-dipeptide derived inhibitors, extension of this work to less peptidyl inhibitors, and the anti-viral properties of these compounds.

As shown in Scheme 1 our synthesis is quite flexible for evaluating structure activity relationships. Diacids 2 are prepared in three steps as described previously ¹ from the appropriate chiral oxazolidinone ³. Treatment of 2 with various amines (R₂NH₂) in the presence of amide coupling reagents afford bis-amides 3. We used dipeptides in our early work for the preliminary evaluation of terminal groups. Bis-amides 3 were converted into alcohols 5 as described previously. Table 1 lists IC₅₀ values as enzyme inhibitors ^{4a} for various dipeptide derived secondary alcohols. Of the derivatives surveyed, those with an isopropyl group in the P₂ position are most potent.

A late branch point in our synthesis was provided by the ease of manipulation of the exocyclic olefin. This allowed us to explore various options. Specifically, oxidation using either ozone or catalytic OsO₄ in the presence of sodium periodate afforded ketones 4. Reduction of the ketone afforded alcohols 5. Catalytic osmylation of 3 afforded diols 6. Treatment of 3 with *m*-chloroperoxybenzoic acid afforded epoxides 7. Table 2 lists IC₅₀ values ^{4a} as enzyme inhibitors for these various derivatives. As expected ¹, secondary alcohols are much more potent than the corresponding ketones, diols and epoxides in a direct comparison. We have not been able to obtain any evidence that epoxide 14 acts as an irreversible enzyme inhibitor.⁵

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Scheme 1

$$R_2$$
 R_2
 R_3
 R_4
 R_5
 R_4
 R_5
 R_5
 R_6
 R_7
 R_8
 R_8
 R_9
 R_9

Table 1 Enzyme inhibiting properties of various dipeptide derived inhibitors of HIV-1 protease.

Compound	P1	P2	Р3	IC50 (nM)4a
8	n-Bu	Me	Me	75
9	n-Bu	<i>i-</i> Bu	Bn	90
10	n-Bu	i-Pr	i-Pr	5
11	Bn	<i>i-</i> Bu	Bn	30
1	Bn	<i>i</i> -Pr	i-Pr	5
12	Bn	i-Pr	Bn	10

Table 2 Enzyme inhibiting properties of various dipeptide derived inhibitors of HIV-1 protease. The effect of central group.

Compound	P1	P2	P3	X	IC50 (nM)4a
9	n-Bu	i-Bu	Bn	Н,ОН	90
13	n-Bu	<i>i</i> -Bu	Bn	OH,CH2OH	2,000
14	<i>n-</i> Bu	<i>i</i> -Bu	Bn	-CH ₂ O-	400
15	Bn	<i>i-</i> Pr	i-Pr	Ō	>10,000

The results of Table 1 and 2 allowed us to define the central unit (4-hydroxy-2,6-bis-(phenylmethyl) heptanediamide) of a pseudo-symmetrical lead structure. At this point there were two divergent strategies for lead optimization: *de novo* design of non-peptidyl end groups ⁶ and traditional medicinal chemistry optimization of 1. Using the data in Table 1 and 2 as well as structure-activity information which was reported in the literature ⁷ we prepared a variety of less peptidyl compounds. Several examples of these compounds are reported in Table 3. Of these compounds, 17 proved to be the most potent inhibitor in each of our enzyme assays.⁴

Table 3 Enzyme inhibiting properties of various monopeptide derived inhibitors of HIV-1 protease.

Compound P₃ P₂
$$IC_{50}$$
 $(nM)^{4a}$ IC_{50} $(nM)^{4b}$ K_{1} $(nM)^{4c}$

16 IC_{50} IC_{50}

Table 4 Ability of compounds to inhibit growth of HIV-1 IIIB

Compound	$IC_{50}(\mu M)$	Compound	$IC_{50}(\mu M)$	Compound	$IC_{50}(\mu M)$
1	1.6	16	2.1	19	>10
11	2.3	17	0.4	20	7.5
12	2.4	18	0.5	Ro 31-89594b	0.003

Many of the compounds reported in Tables 1 and 3 were evaluated for their ability to act as antiviral agents in cell culture. This data is reported in Table 4. Of interest is the lack of a direct correlation between enzyme inhibition (relative potency: 17 > 16 > 18 > 19 > 20) and antiviral activity (relative potency: 17 > 18 > 16 > 20 > 19). The compounds reported in Table 4 have rather high molecular weights, poor solubility properties in all solvents examined, and a poor ratio of antiviral activity to enzyme inhibiting activity. This information led us to speculate that the relatively modest antiviral activity of these compounds was due to low cellular bioavailability as a consequence of their poor physical properties. In addition, compound 17 shows no oral bioavailability in mice.

In an attempt to change the physical properties of 17, unsymmetrical compounds of the general structure 21 were prepared. In 21 "A" represents a group to impart potent enzyme inhibiting properties while "B" represents a group to change physical properties. While these unsymmetrical compounds are expected to be less potent enzyme inhibitors due to lack of P₃ groups, it was hoped that improved physical properties might allow these compounds to have comparable or enhanced antiviral activities.

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In a striking example, compound 22 was prepared and in direct comparison to 17 found to be ca. 8 times less potent ($IC_{50} = 300 \text{ nM}$)^{4b} as an enzyme inhibitor, have significantly improved solubility in several solvents (solubility of 17: 0.02 mg/ml in CHCl₃, < 0.002 mg/ml in H₂O; solubility of 22: 0.5 mg/ml in CHCl₃, 0.02mg/ml in H₂O) and be equipotent ($IC_{50} = 0.55 \mu M$) as an antiviral agent.

The synthesis of 22 is described in Scheme 2. Diacid 2 (R_1 = CH₂Ph) was teated successively with 1.0 mol-equiv. of iso-butyl amine in the presence of the BOP reagent ¹⁰, followed by treatment with an excess of 23 in the presence of BOP. This generated a statistical mixture of symmetrical products 24 and 25 and unsymmetrical product 26. The mixture was separated and 26 was converted in two steps (1. O₃; 2. NaBH₄) into 22 (as an inseparable, 1:1 mixture of diastereomers). Symmetrical olefin 25 had been previously converted into alcohol 27 which was shown to be a weak inhibitor ($IC_{50} > 1000 \text{ nM}$)^{4a} of HIV-1 protease. Other low molecular weight (< 500) symmetrical alcohols, lacking terminal P₃ groups and connecting amides, also proved to be weak inhibitors of the enzyme.

The potent antiviral activity of unsymmetrical compound 22 is consistent with the hypothesis that poor physical properties, of high (> 700) molecular weight symmetrical compounds, manifests itself in less than optimal antiviral activity. This result gives credence to a strategy to prepare compounds of general structure 21, and related strategies, where one part of the molecule is designed to inhibit the enzyme while other part is designed to improve physical properties. Studies which address this strategy will be reported in the future. 11

References and Notes

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- (a) Assay A. These enzyme inhibition studies were performed using HIV-1 protease obtained from Medigenics, Omaha, NE. using a modification of the procedure in Tomaselli, A. G.; Olsen, K. K.; Hui, J. O.; Staples, D. J.; Sawyer, T. K.; Heinrikson, R. L.; Tomich, C-S. C. *Biochem.* 1990, 29, 264-9. As a literature control A-74704 (Erickson, J.; Neidhart, D. J.; VanDrie, J.; Kempf, D. J.; Wang, X. C.; Norbeck, D. W.; Plattner, J. J.;

- Rittenhouse, J. W.; Turon, M.; Wideburg, N.; Kohlbrenner, W. E.; Simmer, R.; Helfrinch, R.; Paul, D. A.; Knigge, M. *Science* 1990, 249, 527-33) was determined to have an $IC_{50} = 25$ nM and Ki = 10 nM (literature report of $IC_{50} = 3$ nM and $K_i = 4.5$ nM).
- (b) Assay B. These enzyme inhibition studies were performed using recombinant HIV-1 protease prepared and purified by Drs. S. Plotch and E. Baum. The enzyme assay was an ¹²⁵I-SPA (scintillation proximity assay) using a kit provided by Amersham International plc, Bucks, England. Using the procedure described by the manufacturer, enzyme and test compounds were incubated with the bead suspension for 5 min., following termination of the reaction the radioactivity of the assay mixture was determined. Enzyme inhibition was determined as a function of enzyme concentration. As a literature control Ro 31-8959 (Roberts, N. A.; Martin, J. A.; Kinchington, D.; Broadhurst, A. V.; Craig, J. C.; Duncan, I. B.; Galpin, S. A.; Handa, B. K.; Kay, J.; Krohn, A.; Lambert, R. W.; Merrett, J. H.; Mills, J.S.; Parkes, K. E. B.; Redshaw, S.; Ritchie, A. J.; Taylor, D. L.; Thomas, G. L.; Machin, P. J. Science 1990, 248, 358-61) was determined to have an IC₅₀ = 30 nM and Ki = 3 nM (literature report of IC₅₀ < 0.4 nM).
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