

Cyclic HIV-1 Protease Inhibitors Derived from Mannitol: Synthesis, Inhibitory Potencies, and Computational Predictions of Binding Affinities

Johan Hultén,[†] Nicholas M. Bonham,[†] Ulrika Nillroth,[‡] Tomas Hansson,[§] Guido Zuccarello,^{||} Abderrahim Bouzide,^{||} Johan Åqvist,[§] Björn Classon,[∇] U. Helena Danielson,[‡] Anders Karlén,[†] Ingemar Kvarnström,^{||} Bertil Samuelsson,[⊥] and Anders Hallberg^{*,†}

Department of Organic Pharmaceutical Chemistry, Uppsala Biomedical Center, Uppsala University, Box 574, S-751 23 Uppsala, Sweden, Department of Biochemistry, Uppsala Biomedical Center, Uppsala University, Box 576, S-751 23 Uppsala, Sweden, Department of Molecular Biology, Uppsala Biomedical Center, Uppsala University, Box 590, S-751 24 Uppsala, Sweden, Department of Chemistry, Linköping University, S-581 83 Linköping, Sweden, Medivir AB, Lunastigen 7, S-141 44 Huddinge, Sweden, and Department of Organic Chemistry, Arrhenius Laboratory, Stockholm University, S-106 91 Stockholm, Sweden

Received October 17, 1996[⊗]

Ten C_2 -symmetric cyclic urea and sulfamide derivatives have been synthesized from L-mannonic γ -lactone and D-mannitol. The results of experimental measurement of their inhibitory potencies against HIV-1 protease were compared to calculated free energies of binding derived from molecular dynamics (MD) simulations. The compounds were selected, firstly, to enable elucidation of the role of stereochemistry for binding affinity (**1a–d**) and, secondly, to allow evaluation of the effects of variation in the link to the P1 and P1' phenyl groups on affinity (**1a** and **2–5**). Thirdly, compounds with hydrogen bond-accepting or -donating groups attached to the phenyl groups in the P2 and P2' side chains (**6** and **7**) were selected. Binding free energies were estimated by a linear response method, whose predictive power for estimating binding affinities from MD simulations was demonstrated.

Introduction

Human immunodeficiency virus (HIV) is the causative agent of acquired immune deficiency syndrome, AIDS.¹ HIV protease processes the viral polyproteins Pr55_{gag} and Pr160_{gag-pol} into structural proteins and enzymes, including the protease itself. The activity of the protease is vital for proper assembly and maturation of infectious virions.² Therefore HIV protease is an important chemotherapeutic target.³

Numerous examples of potent inhibitors of HIV-1 protease have been disclosed.⁴ A common feature observed in the X-ray crystal structures of linear inhibitor–HIV-1 protease complexes is the presence of a tetracoordinated structural water molecule linking the bound inhibitors to the flexible flaps of the HIV-1 protease dimer.⁵ On the basis of this structural information, Lam et al. have performed an elegant design of nonpeptide cyclic ureas, constituting an entirely new class of HIV-1 protease inhibitors.⁶ More recently, Sham et al. have reported a series of new azacyclic ureas possessing high potency as inhibitors of HIV-1 protease.⁷ The fundamental characteristic of these inhibitors, e.g., DMP 323 and A-98881 (Figure 1), is the cyclic urea carbonyl oxygen mimicking the structural water molecule.⁶

The application of computer modeling and simulation of molecular interactions to drug design is a field currently in rapid growth. Methods allowing reliable predictions of affinity and specificity of ligands would be of paramount value in the discovery process, aiding

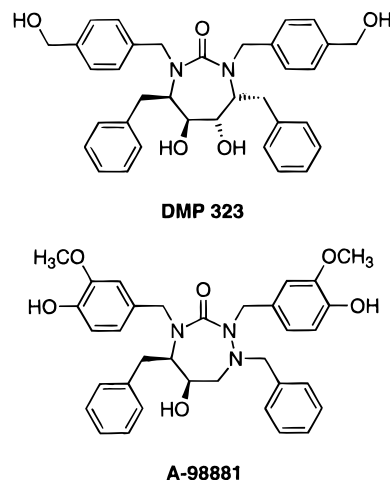


Figure 1. Structures of DMP 323 and A-98881.

in the selection of ligands with high potential before synthesis is commenced. In cases where the three-dimensional structure of the drug target is known, methods for scoring putative inhibitors from single conformations of ligand–target complexes have been developed.⁸ But even with a 3D-structure of the target available, computational predictions of binding affinity for new ligands is difficult in the absence of accurate 3D-structures of the corresponding complexes. The main reason for this lies in the difficulty of predicting the exact structure of the complex. Force field-based structural optimization and scoring has been applied recently to HIV-1 protease inhibitors with promising results.⁹ Åqvist et al. have described another approach, which has yielded good results for several systems,¹⁰ in which absolute binding free energies are estimated by a linear response approximation from thermal averaging of the structure and interaction energies by molec-

* Corresponding author.

[†] Department of Organic Pharmaceutical Chemistry.

[‡] Department of Biochemistry.

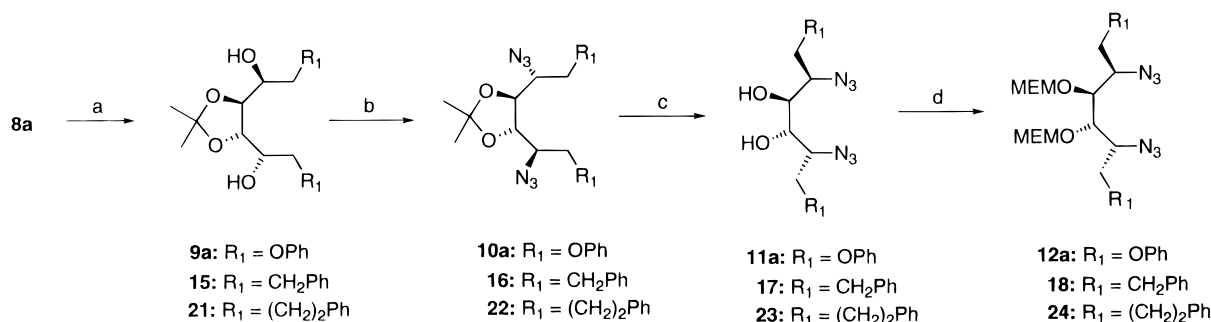
[§] Department of Molecular Biology.

^{||} Linköping University.

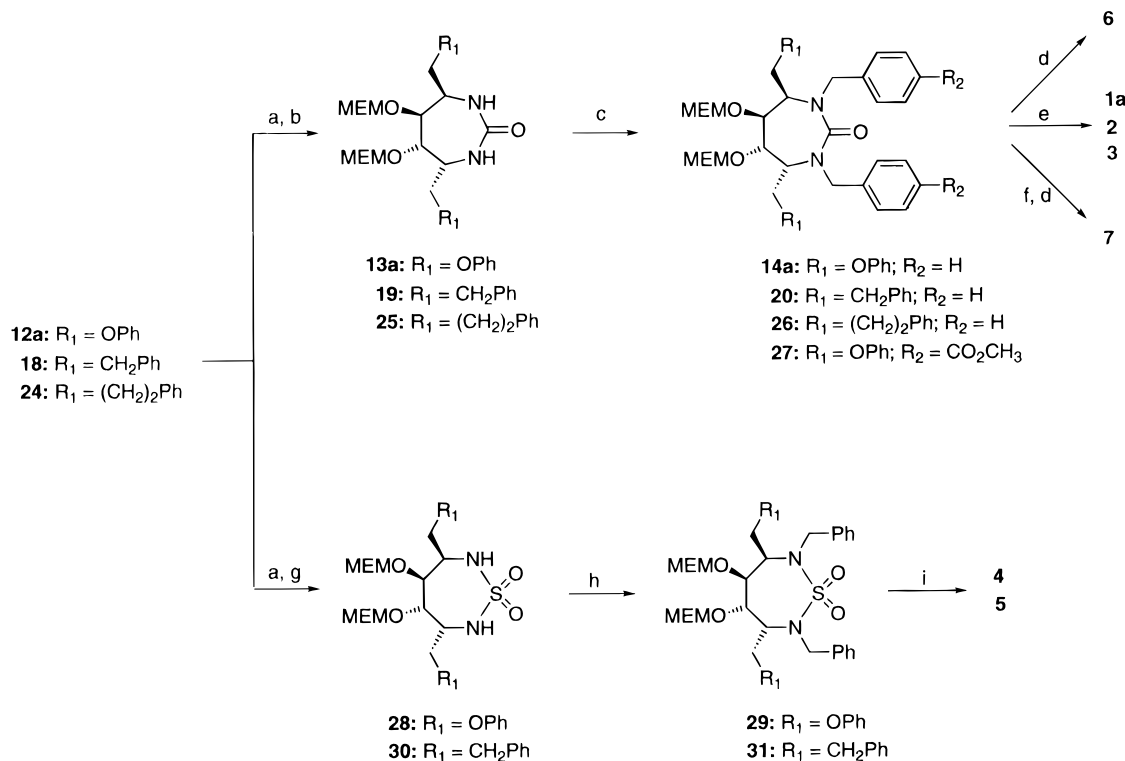
[∇] Medivir AB.

[⊥] Stockholm University.

[⊗] Abstract published in *Advance ACS Abstracts*, December 15, 1996.

Scheme 1^a

^a (a) PhOH, NaH, THF or BnMgCl, CuI, THF or Ph(CH₂)₂Br, Mg, CuI, THF; (b) DEAD, DPPA, PPh₃; (c) CH₃CN, 3 M HCl; (d) MEMCl, NaH, THF.

Scheme 2^a

^a (a) H₂, Pd/C (10%), EtOAc; (b) CDI, CH₂Cl₂; (c) BnBr or methyl 4-(bromomethyl)benzoate, NaH, DMF; (d) CH₃OH, concentrated HCl; (e) CH₃CN, 3 M HCl; (f) LiBH₄, ether; (g) sulfamide, pyridine; (h) BnBr, NaH, DMF; (i) HCl in ether, methanol.

ular dynamics (MD) simulations. This procedure was applied recently to DMP 323 and two linear peptide HIV-1 protease inhibitors.¹¹

Several groups have used carbohydrates as chiral precursors for the synthesis of linear HIV-1 protease inhibitors.¹² Here we describe the synthesis of 10 cyclic C₂-symmetric compounds related to DMP 323, which were prepared from the readily available L-mannonic γ -lactone or D-mannitol.¹³ For these 10 model compounds, we compare our results from experimental elucidation of their inhibitory potencies against HIV-1 protease to calculated free energies of binding derived from MD simulations. The compounds were selected to enable comparison of (a) the C₂-symmetric diastereomers **1a–d**, (b) the ureas **1a**, **2**, and **3** and the sulfamides **4** and **5**, which were modified in the link to the P1 and P1' phenyl groups, and (c) the compounds **6** and **7**, modified in the P2 and P2' side chains with groups combining hydrogen bond acceptor and donor capacity, respectively.

Results

Chemistry. The diepoxides **8a,c** were used as key intermediates for the synthesis of **1a**, **2–7**, and **1c**, respectively, and the diepoxides **8b,d** for the synthesis of **1b,d**, respectively (Figure 3). For the preparation of the diepoxides **8b,d** D-mannitol was used as starting material.¹⁴ We employed this procedure for the preparation of **8a,c** from L-mannonic γ -lactone.¹³ The synthetic methods used for the preparation of **1a–d** and **2–7** are shown in Tables 1–4, and the syntheses of the *RSSR* stereoisomers **1a** and **2–7** are outlined in Schemes 1 and 2.

Nucleophilic ring opening of the diepoxides occurs with high regio- and stereoselectivity,¹⁵ and accordingly, ring opening of **8a**, shown in Scheme 1, and **8b** by phenolate smoothly provided **9a,b**^{15c} after 7 h. Ring opening of **8c,d** provided the corresponding diastereomers **9c,d** after 24 h reaction times, in yields ranging of 61–72%. Standard Mitsunobu reaction¹⁶ lead to the azides **10a–d**, with inverted configuration at carbons

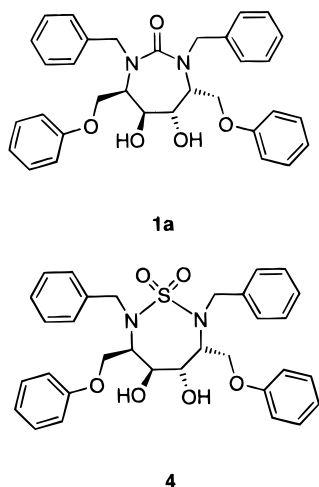


Figure 2. Structures of one urea (**1a**) and one sulfamide (**4**) compound with *RSSR* stereochemistry.

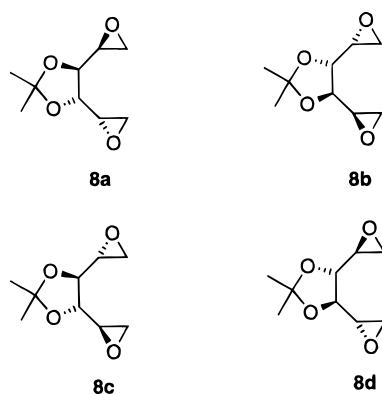


Figure 3. Structures of the bisepoxides **8a-d**.

2 and 5. Reduction to the primary amines occurred smoothly, but cyclization was not facile, probably due to ring strain imposed by the *trans*-fused five-membered acetonide ring.¹⁷ Deketalization of the azides furnished **11a-d**. Treatment with MEM-chloride¹⁸ afforded the bis-MEM ethers **12a-d** in 83–88% isolated yields. Hydrogenation on Pd/C provided the crude amines, which were reacted with carbonyldiimidazole¹⁹ to give the cyclic ureas **13a-d** in 55–68% yields from the appropriate azide. Benzylation²⁰ to give **14a** and the enantiomer **14b** was complete in 12 h, while formation of **14c,d** required 24 h reaction time for full conversion.²¹ Deprotection gave the target compounds **1a-d** in 6–17% overall yield from **8a-d**.

For the preparation of **2**¹³ and **3** a similar reaction sequence was adopted. Ring opening of **8a** with benzylmagnesium chloride or phenylethylmagnesium bromide in the presence of CuI^{2a} delivered **15** and **21**, respectively. The Mitsunobu reaction followed by de-ketalization and re-protection with MEM-chloride provided **18** and **24**. After hydrogenation, cyclization to give the ureas **19** and **25** occurred smoothly. Benzylation and subsequent deprotection furnished **2** and **3** in 14–18% overall yield from the epoxide. Alkylation with methyl 4-(bromomethyl)benzoate and deprotection provided the ester **6**, while reduction²² before the deprotection furnished the hydroxymethyl derivative **7**. Sulfamides **4** and **5**^{6b} were prepared by heating the crude amine with sulfamide in refluxing pyridine. This cyclization²³ followed by benzylation using sodium hydride as base in DMF and subsequent deprotection with dry

Table 1. Structures and Preparation Methods for Mannitol Derivatives

compd ^a	config	R ¹	X	prepn method ^b
9a	<i>SSSS</i>	OPh	OH	I
9b	<i>RRRR</i>	OPh	OH	I
9c	<i>RSSR</i>	OPh	OH	I
9d	<i>SRRS</i>	OPh	OH	I
10a	<i>RSSR</i>	OPh	N ₃	II
10b	<i>SRRS</i>	OPh	N ₃	II
10c	<i>SSSS</i>	OPh	N ₃	II
10d	<i>RRRR</i>	OPh	N ₃	II
15	<i>SSSS</i>	CH ₂ Ph	OH	VIII
16	<i>RSSR</i>	CH ₂ Ph	N ₃	II
21	<i>SSSS</i>	(CH ₂) ₂ Ph	OH	IX
22	<i>RSSR</i>	(CH ₂) ₂ Ph	N ₃	II

^a For experimental data of compounds **9b-d** and **10b-d**, see Supporting Information. ^b See the Experimental Section.

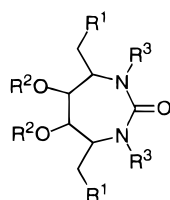
Table 2. Structures and Preparation Methods for Mannitol Derivatives

compd ^a	config	R ¹	R ²	X	prepn method ^b
11a	<i>RSSR</i>	OPh	H	N ₃	III
11b	<i>SRRS</i>	OPh	H	N ₃	III
11c	<i>SSSS</i>	OPh	H	N ₃	III
11d	<i>RRRR</i>	OPh	H	N ₃	III
12a	<i>RSSR</i>	OPh	MEM	N ₃	IV
12b	<i>SRRS</i>	OPh	MEM	N ₃	IV
12c	<i>SSSS</i>	OPh	MEM	N ₃	IV
12d	<i>RRRR</i>	OPh	MEM	N ₃	IV
17	<i>RSSR</i>	CH ₂ Ph	H	N ₃	III
18	<i>RSSR</i>	CH ₂ Ph	MEM	N ₃	IV
23	<i>RSSR</i>	(CH ₂) ₂ Ph	H	N ₃	III
24	<i>RSSR</i>	(CH ₂) ₂ Ph	MEM	N ₃	IV

^a For experimental data of compounds **11b-d** and **12b-d**, see Supporting Information. ^b See the Experimental Section.

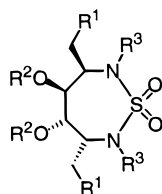
HCl in ether–methanol afforded compounds **4** and **5** in 18% and 16% overall yield from the diepoxide as depicted in Scheme 2.

Inhibition of HIV-1 Protease. The inhibitory effect of the synthesized compounds was determined with purified HIV-1 protease in a standardized assay.²⁴ The results are presented as IC₅₀-values, i.e., the concentration of inhibitor resulting in 50% inhibition in this assay (Table 5). Since some compounds were not sufficiently potent to allow determination of IC₅₀-values, results are also given as percent inhibition at 10 μM (Table 5). Compounds that showed significant inhibition were further characterized, and K_i-values were determined (Table 5). The model for tight-binding inhibitors used for analysis²⁵ was adequate except for the most potent inhibitors, as judged by the correlation between k_{cat} and K_m-values, obtained when fitting data for different inhibitors and by statistical criteria (not shown). When the K_i-values are lower than the enzyme concentration used, the portion of free inhibitor becomes insignificant, since the enzyme is essentially titrated with inhibitor under these conditions; thus, in such cases the method

Table 3. Structures and Preparation Methods for Cyclic Urea Compounds

compd ^a	config	R ¹	R ²	R ³	prepn method ^b
1a	<i>RSSR</i>	OPh	H	CH ₂ Ph	VII
1b	<i>SRRS</i>	OPh	H	CH ₂ Ph	VII
1c	<i>SSSS</i>	OPh	H	CH ₂ Ph	VII
1d	<i>RRRR</i>	OPh	H	CH ₂ Ph	VII
2	<i>RSSR</i>	CH ₂ Ph	H	CH ₂ Ph	VII
3	<i>RSSR</i>	(CH ₂) ₂ Ph	H	CH ₂ Ph	VII
6	<i>RSSR</i>	OPh	H	CH ₂ -4-(CO ₂ CH ₃)Ph	X
7	<i>RSSR</i>	OPh	H	CH ₂ -4-(CH ₂ OH)Ph	XI
13a	<i>RSSR</i>	OPh	MEM	H	V
13b	<i>SRRS</i>	OPh	MEM	H	V
13c	<i>SSSS</i>	OPh	MEM	H	V
13d	<i>RRRR</i>	OPh	MEM	H	V
14a	<i>RSSR</i>	OPh	H	CH ₂ Ph	VI
14b	<i>SRRS</i>	OPh	MEM	CH ₂ Ph	VI
14c	<i>SSSS</i>	OPh	MEM	CH ₂ Ph	VI
14d	<i>RRRR</i>	OPh	MEM	CH ₂ Ph	VI
19^c	<i>RSSR</i>	CH ₂ Ph	MEM	H	V
20	<i>RSSR</i>	CH ₂ Ph	MEM	CH ₂ Ph	VI
25^c	<i>RSSR</i>	(CH ₂) ₂ Ph	MEM	H	V
26	<i>RSSR</i>	(CH ₂) ₂ Ph	MEM	CH ₂ Ph	VI
27^d	<i>RSSR</i>	OPh	MEM	CH ₂ -4-(CO ₂ CH ₃)Ph	VI

^a For experimental data of compounds **13b–d** and **14b–d**, see Supporting Information. ^b See the Experimental Section. ^c 2 days reaction time. ^d Methyl 4-(bromomethyl)benzoate was used instead of BnBr.

Table 4. Structures and Preparation Methods for Cyclic Sulfamide Compounds

compd	config	R ¹	R ²	R ³	prepn method ^a
4	<i>RSSR</i>	OPh	H	CH ₂ Ph	XIV
5	<i>RSSR</i>	CH ₂ Ph	H	CH ₂ Ph	XIV
28	<i>RSSR</i>	OPh	MEM	H	XII
29	<i>RSSR</i>	OPh	MEM	CH ₂ Ph	XIII
30	<i>RSSR</i>	CH ₂ Ph	MEM	H	XII
31	<i>RSSR</i>	CH ₂ Ph	MEM	CH ₂ Ph	XIII

^a See the Experimental Section.

Table 5. Inhibitory Activity of Cyclic C₂-Symmetric Urea and Sulfamide Compounds against HIV-1 Protease

compd	config	inhibition (%) (at 10 μM)	IC ₅₀ (μM)	K _i (nM)		ΔG _{bind} (kcal/mol)	
				expt ^a	calcd	expt	calcd
DMP 323	<i>RSSR</i>	nd	0.015	<1 ^b	0.083	<-12.8	-14.3 ± 1.0
1a	<i>RSSR</i>	nd	0.08	12.2	1.1	-11.2	-12.7 ± 0.5
1b	<i>SRRS</i>	15	nd	nd	15		-11.1 ± 0.7
1c	<i>SSSS</i>	10	nd	nd	3200		-7.8 ± 0.9
1d	<i>RRRR</i>	0	nd	nd	36000		-6.3 ± 0.5
2	<i>RSSR</i>	nd	4.0	214	7.8	-9.5	-11.5 ± 1.4
3	<i>RSSR</i>	nd	10.0	570	0.80	-8.9	-12.9 ± 0.7
4	<i>RSSR</i>	nd	0.200	19.1	0.80	-10.9	-12.9 ± 1.7
5	<i>RSSR</i>	nd	0.200	14.7	0.94	-11.1	-12.8 ± 0.5
6	<i>RSSR</i>	nd	8.0	nd	9.2		-11.4 ± 0.9
7	<i>RSSR</i>	nd	0.010	<1 ^b	0.051	<-12.8	-14.6 ± 1.6

^a The standard error was less than 10%. ^b K_i(DMP 323) = 0.27 nM and K_i(**7**) = 0.23 nM have been determined by a different method (Nillroth et al., to be published). nd = not determined.

cannot be used. The standard errors for determination of the parameters were ≤10%.

The stereochemistry of the compounds was found to be critical for their inhibitory effect. Only the *RSSR* isomer **1a** inhibits the enzyme significantly, K_i = 12.2 nM. The other isomers (**1b–d**) showed little (<15%) or no inhibitory effect even at micromolar concentrations. Small structural modifications of compound **1a** gave significant changes in the inhibitory potency. Changing the P1/P1' side chains from phenoxyethyl (**1a**) to phenylethyl (**2**) resulted in a 20-fold reduction of the K_i-value. A 50-fold loss of inhibitory effect relative to **1a** was found when the P1/P1' side chain was further extended to give the phenylpropyl derivative **3**. The cyclic sulfamides **4** and **5** both showed inhibitory potencies of the same order of magnitude as **1a**. Interestingly, exchanging the P1/P1' ether oxygens for methylene groups in the sulfamides gave essentially equipotent derivatives (**4** vs **5**), contrary to the observation with the ureas (**1a** vs **2**). Finally, both of the *p*-hydroxymethyl compounds (**7** and DMP 323) exhibited high potencies, while compound **6** exhibited lower binding affinity than **1a**.

Dynamics Simulations and Free Energy Calculations. Binding free energies were estimated by an approximation described elsewhere,¹⁰ which we will refer to here as the linear interaction energy (LIE) method. Two MD simulations were performed for each inhibitor, one for the inhibitor free in water and the other for the inhibitor bound inside the solvated protein. The simulation results are summarized in Table 6 in terms of the average electrostatic (polar) and van der Waals (nonpolar) ligand interaction energies for the bound and unbound states. Hydrogen bonds are included implicitly in these two quantities and are mainly reflected in the electrostatic energy. Table 6 also gives the resulting free energies of binding obtained with the LIE approximation. The predicted absolute binding free energies are in reasonable agreement with the experimental values (Table 5) but are generally ca. 2 kcal/mol too negative. The simulations identify **1a** as a good inhibitor and give the correct ranking of the compounds **1a–d**. In particular, the two compounds **1c,d** are found to bind poorly, while the binding energy of **1b** is somewhat overestimated. The water simulations of the enantiomeric pairs converged as expected to give very similar values within each pair. The average of the two simulations is given in Table 6 for each pair. For the compounds **2** and **3**, with the ether oxygens replaced by methylene and ethylene groups, **2** is predicted to be

Table 6. Dynamics Averages of Nonbonded Terms in the Simulations^a

compd	$\langle V_{\text{vdw}} \rangle_{\text{prot}}$	$\langle V_{\text{vdw}} \rangle_{\text{wat}}$	$\langle V_{\text{el}} \rangle_{\text{prot}}$	$\langle V_{\text{el}} \rangle_{\text{wat}}$	ΔG_{bind}		
					vdw	el	calcd
DMP 323	-86.57 ± 1.17	-42.07 ± 0.19	-82.22 ± 1.08	-67.89 ± 0.48	-7.16 ± 0.2	-7.16 ± 0.8	-14.3 ± 1.0
1a	-85.76 ± 0.17	-46.66 ± 0.88	-50.99 ± 0.10	-38.19 ± 0.44	-6.30 ± 0.2	-6.40 ± 0.3	-12.7 ± 0.5
1b	-90.61 ± 1.08	-46.66 ± 0.88	-46.31 ± 0.20	-38.19 ± 0.44	-7.08 ± 0.3	-4.06 ± 0.4	-11.1 ± 0.7
1c	-88.30 ± 0.70	-46.63 ± 0.12	-41.42 ± 1.30	-39.32 ± 0.12	-6.71 ± 0.2	-1.05 ± 0.7	-7.8 ± 0.9
1d	-86.79 ± 0.52	-46.63 ± 0.12	-38.99 ± 0.65	-39.32 ± 0.12	-6.47 ± 0.1	+0.16 ± 0.4	-6.3 ± 0.5
2	-86.32 ± 0.21	-48.42 ± 0.25	-45.17 ± 1.66	-34.29 ± 0.97	-6.10 ± 0.1	-5.44 ± 1.3	-11.5 ± 1.4
3	-90.37 ± 0.23	-52.59 ± 0.44	-50.57 ± 0.21	-37.00 ± 0.96	-6.08 ± 0.1	-6.78 ± 0.6	-12.9 ± 0.7
4	-88.85 ± 0.03	-50.91 ± 0.09	-49.70 ± 1.91	-36.09 ± 1.31	-6.11 ± 0.1	-6.80 ± 1.6	-12.9 ± 1.7
5	-88.42 ± 0.04	-49.11 ± 0.04	-48.97 ± 0.22	-36.01 ± 0.57	-6.33 ± 0.1	-6.48 ± 0.4	-12.8 ± 0.5
6	-103.61 ± 0.10	-53.65 ± 1.39	-57.38 ± 0.63	-50.68 ± 0.51	-8.04 ± 0.3	-3.35 ± 0.6	-11.4 ± 0.9
7	-90.86 ± 0.43	-44.13 ± 0.53	-84.98 ± 1.33	-70.78 ± 1.34	-7.52 ± 0.2	-7.10 ± 1.4	-14.6 ± 1.6

^a Values are averages over 125 ps (250 ps for DMP 323 and 7). The error estimates are one-half the differences between the first and last 62.5 ps (or 125 ps) and thus measure precision rather than accuracy. All values are given in kcal/mol. The contributions from the two terms in the binding free energy approximation are also shown.

less active than **1a**, in agreement with the binding experiments. However, the calculated binding energy for **3** is about the same as for **1a**, which is a significant overestimation compared to experimental values. Calculations on the sulfamides **4** and **5** yielded binding free energies close to those obtained for **1a**, in agreement with experimental findings. The ester **6** was predicted to be similarly potent to **2**, which is indeed also indicated by the experimentally observed IC₅₀-value. The calculations predict essentially equal, large negative, binding energies for DMP 323 and **7**, pointing to a negligible effect of the added oxygen. There are, however, significant uncertainties in this prediction, since we find that a major part of the interaction energy of these two inhibitors arises from interactions of the hydroxymethyl groups with aspartates 30/230, which are mobile and in contact with bulk water, leading to large variations during simulation time. For these two inhibitors, data collection times for the protein simulations were therefore doubled. The free energy of binding obtained for DMP 323 is more negative than has been reported earlier¹¹ due to the interaction with Asp 30/230, since the aspartate side chains need to move (χ_1 rotation) from their crystallographic starting positions for this interaction to occur, a conformational change that was neither anticipated nor observed in the calculations on DMP 323.¹¹ As noted above, exact experimental values of the inhibition constants could not be obtained for the potent inhibitors **7** and DMP 323 (K_i of 0.27 nM has been reported for DMP 323).⁶

The MD average structures from the calculations give a picture of binding arrangements over some tens of picoseconds. On this time scale, all inhibitors were found to bind somewhat asymmetrically in the binding pocket, especially **1a,b**. The reason for this asymmetry of binding was the protonation state chosen for the catalytic aspartates. Asp 25 was considered to be negatively charged, Asp 225 was protonated,²⁶ and both hydroxyls of the inhibitor were found to point toward Asp 25, tilting the seven-membered ring slightly. Strong hydrogen bonds can form between diequatorial hydroxyl groups and the charged aspartate. Only one of the diaxial hydroxyl groups found in relaxed conformations of the free compounds **1c,d** would be able to hydrogen bond to Asp 25. A strong charge-dipole interaction would then be replaced by a weaker dipole-dipole interaction with Asp 225, leading to weaker binding. For **1c**, the calculations indeed predict a conformational change in the seven-membered cyclic urea ring, yielding

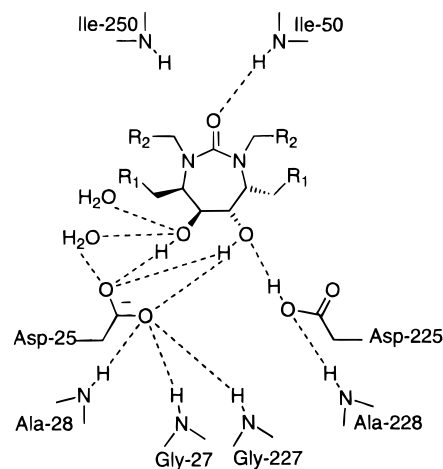


Figure 4. Schematic picture showing the general hydrogen-bonding arrangements in the vicinity of the inhibitors with *RSSR* stereochemistry, on the tens-of-picoseconds time scale of the simulations. Distances in the figure are not to scale, and all water molecules depicted are in contact with bulk water. Dashed lines indicate that atom distances were on average favorable for hydrogen bond formation.

diequatorial hydroxyls, but this seems to be at the expense of other favorable interactions and still results in poor binding. The average structures for the *RSSR* inhibitors are summarized in Figure 4, which summarizes the average hydrogen-bonding arrangement in the vicinity of the inhibitor on the time scale of the simulations.

Discussion

High binding affinity is attained by both steric and electrostatic complementarity between the ligand and its target. Imperfect steric fit generally can be relaxed by small adjustments in the protein. Electrostatic complementarity, which involves both polar interactions (e.g., hydrogen bonds) and nonpolar interactions (e.g., hydrophobic effects), usually cannot be adjusted easily due to the long-range nature of electrostatic fields and is therefore a major determinant of binding specificity. These effects are apparent in the results for the stereoisomers **1a-d**. For all four compounds, shape complementarity alone provides a basic level of affinity, as demonstrated by the van der Waals term of the MD simulation results, but the exact polar-hydrophobic matching within this shape results in the observed (Table 5) large differences, leading to high affinity in

some cases (in the nanomolar range for **1a**). The present results indicate that the stereochemistry of the compounds is critical for their inhibitory potency. It is especially important for the hydroxyl groups to be positioned optimally, and the *RSSR* stereoconfiguration of **1a** enables favorable hydrogen bonds to the charged aspartate.

The MD calculations suggest the following explanation for the 20-fold enhancement of binding affinity of **1a** as compared **2**: A dipole is formed by each partially negative ether oxygen and its two partially positive neighboring carbons, since the angle between these three atoms is less than 180°. Analysis of the average structure from the MD simulations shows that for **1a**, these dipoles are aligned to an electrostatic field arising from the protein environment, with the major contribution from the charged Asp 25 but also with significant contributions from the flap peptide bonds from residues 49–50 and 249–250.

The cyclic sulfamides **4** and **5** were synthesized with *RSSR* stereochemistry since this configuration was found to be optimal for the urea compounds. The binding energies measured for the sulfamide compounds **4** and **5** are both approximately equal to that found for **1a**, in agreement with the result derived from the MD simulations. One might expect the same electrostatic effects in **4** and **5** as in **1a** and **2**, but the average structures of **4** and **5** predict that there is not enough space available for the ether oxygens to be aligned to the field, due to the extra volume from the sulfonyl group compared to the carbonyl group. Thus, no decrease in affinity results from the substitution of the ether oxygen in compound **4** with the methylene group in compound **5**.

Compounds **6** and **7** were synthesized in order to study hydrogen-bonding capacity in the S2/S2' cavity. As expected, **7** exhibited high binding affinity in analogy to DMP 323. This is probably due to the ability of **7** to interact with the negatively charged aspartates 30/230, as discussed above. The simulations point to a large net loss of electrostatic binding energy of **6** compared to **1a**. This illustrates a general principle of ligand binding energetics, namely, that a comparison always must be made to the unbound reference state, where the ligand is free in water. Therefore, not only must the addition of a group to the ligand lead to favorable interactions in the bound state, but the interactions with the receptor must be stronger than the interactions of the same added group with surrounding water molecules of the unbound state, or no net increase in binding energy will be achieved (Table 6).

One interesting structural feature emerging from the simulations is that the hydrogen bonds between the inhibitor and the flaps seem not to be as strong as possible, as reflected by the fact that only one strong hydrogen bond to the NH groups 50/250 is observed in the tens-of-picoseconds average structure (Figure 4). This could indicate that the cyclic urea carbonyl group might not be the optimal mimic of the crystallographically found water molecule, but this could also be due to imperfections in the representation of the groups involved in the force field used in this work. Moreover it is interesting to note that water molecules (in contact with bulk water) can interact with the cyclic urea ring hydroxyls even when the inhibitor is bound to the protein.

The asymmetric protonation state²⁶ of the catalytic Asp 25/225 used in the present simulations seems to explain the asymmetric binding mode of the inhibitors on the short time scales of the simulations, depicted in Figure 4. However, even if this asymmetry is correct on the short time scale of the simulations, it need not be observable necessarily in crystallographic studies, since these involve implicit averaging over many protein dimers that need not be isomorphous with respect to the aspartate protonation throughout the crystal. Instead, such experiments may reflect the average of Figure 4 and its symmetry-related counterpart, in addition to any contributions from other protonation states, depending on the pH of the crystal.

Conclusion

Synthesis of these inhibitors from readily available L-mannonic γ -lactone or D-mannitol is relatively straightforward, and the ether oxygens, easily incorporated by the synthetic methodology employed, in fact seem to enhance binding 20-fold as compared to the corresponding methylene analog (**2**) in the cyclic urea series. Furthermore, the methylene analog in the sulfamide series (**5**) is 15-fold more potent than the corresponding cyclic urea derivative, but steric effects seem to preclude addition of these two enhancement factors.

The predictive power of the LIE method for estimating binding affinities for the ligands from MD simulations has been demonstrated. The enzymatic measurements give the following ranking of inhibitors (in decreasing potency) DMP 323, **7** > **1a**, **4**, **5** > **2**, **3**, **6** > **1b–d**, whereas the theoretical predictions are DMP 323, **7** > **1a**, **3–5** > **1b**, **2**, **6** > **1c,d**, showing that the relative ordering of the inhibitors is well reproduced by the calculations, with the exception of the large overestimation of the potency of compound **3** and the slight overestimation for compound **1b**.

The reason for the apparently systematic overestimation of the binding strength of the compounds studied here by ca. 2 kcal/mol is not clear. It is, of course, conceivable that our linear response formula with its current parameterization is not accurate enough (although earlier calculations suggested so). Another possibility is that differences between the experimental conditions and the simulation model (e.g., ionic strength) affect the result in a systematic way. A third possibility is that some force field deficiency is responsible for the error. One may, for example, question the non-polar-extended atom representation of phenyl groups in the GROMOS force field²⁷ in this context. This somewhat oversimplified model could lead to an exaggeration of the hydrophobic effect which may be especially significant here since the compounds studied each have four phenyl groups, in contrast to those considered in earlier applications of the method. Nevertheless, the results obtained here suggest that the LIE method is useful as a tool in design projects where the structure of the molecular target is known.

Experimental Section

Chemistry. General Information. Melting points were recorded on an Electrothermal melting point apparatus and are uncorrected. Optical rotations were obtained on a Perkin-Elmer 241 polarimeter. Specific rotations ($[\alpha]_D$) are reported in deg/dm, and the concentration (*c*) is given in g/100 mL in the specific solvent. ¹H and ¹³C NMR spectra were recorded

on a JEOL JNM-EX 270 spectrometer at 270.2 and 67.8 MHz, respectively, or on a JEOL JNM-EX 400 spectrometer at 399.8 and 100.5 MHz, respectively; the chemical shifts are given in ppm relative to tetramethylsilane. Infrared spectra were recorded on a Perkin-Elmer 1600 series FTIR instrument. Elemental analyses were performed by Mikro Kemi AB, Sweden, or Analytische Laboratorien, Germany, and were within $\pm 0.4\%$ of calculated values. Mass spectroscopy was carried out on a JEOL SX 102 instrument. Flash column chromatography was performed on silica gel 60, 0.04–0.063 mm (E. Merck), with gradient elution unless otherwise noted. Thin-layer chromatography was performed on precoated silica gel F-254 plates (0.25 mm; E. Merck) and visualized with UV light and H_2SO_4 in ethanol, phosphomolybdic acid, or ninhydrin. Standard workup: organic layers were dried with MgSO_4 and concentrated *in vacuo*.

3,4-O-Isopropylidene-1,2,5,6-dianhydro-D-iditol (8c): mp 69–70 °C; IR (KBr) ν 2994, 1379, 1249, 1180 cm^{-1} ; $[\alpha]_{\text{D}} = +17.5^\circ$ ($c = 1.20$, CHCl_3 , 24 °C); $^1\text{H NMR}$ (270.2 MHz, CDCl_3) δ 3.85 (dd, $J = 3.1, 1.6$ Hz, 2H, CHOC), 3.07 (m, 2H, CHO), 2.84 (dd, $J = 5.1, 4.1$ Hz, 2H, CH_2O), 2.74 (dd, $J = 5.1, 2.1$ Hz, 2H, CH_2O), 1.40 (s, 6H, CH_3); $^{13}\text{C NMR}$ (67.8 MHz, CDCl_3) δ 110.8, 78.1, 51.2, 43.9, 26.8. Anal. ($\text{C}_9\text{H}_{14}\text{O}_4$) C, H.

3,4-O-Isopropylidene-1,6-di-O-phenyl-L-mannitol (9a). **Method I.** To phenol (8.6 g, 91.5 mmol) in THF/toluene (1:3, 500 mL) was added NaH (1.8 g, 60.8 mmol) under N_2 atmosphere; this was stirred for 20 min at 25 °C. The diepoxide **8a** (2.83 g, 15.2 mmol) was added, and the temperature was raised to 95 °C for 7 h. The solution was allowed to cool and washed with 1 M NaOH (2×150 mL). The combined aqueous extracts were extracted with ether (100 mL). The combined organic extracts were dried and concentrated. Purification by flash chromatography ($\text{CH}_2\text{Cl}_2/\text{CH}_3\text{OH}$, 100:1) gave the product as a white solid (4.09 g, 72%): mp 108–111 °C; IR (CH_2Cl_2) ν 3600–3300, 3050, 2932, 1594, 1492 cm^{-1} ; $[\alpha]_{\text{D}} = -38.0^\circ$ ($c = 1.03$, CHCl_3 , 25 °C); $^1\text{H NMR}$ (270.2 MHz, CDCl_3) δ 7.3 (dd, $J = 7.5, 6.9$ Hz, 4H, OArH[m]), 6.95 (m, 6H, OArH[o + p]), 4.27 (d, $J = 7.6$ Hz, 2H, CHOC), 4.05 (m, 6H, CHOH , CH_2OPh), 3.6 (s, 2H, OH), 1.39 (s, 6H, CH_3); $^{13}\text{C NMR}$ (67.8 MHz, CDCl_3) δ 158.5, 129.4, 121.1, 114.7, 109.8, 79.7, 71.7, 69.4, 26.9. Anal. ($\text{C}_{21}\text{H}_{26}\text{O}_6$) C, H.

2,5-Diazido-3,4-O-isopropylidene-1,6-di-O-phenyl-2,5-dideoxy-D-iditol (10a). **Method II.** To a solution of compound **9a** (0.98 g, 2.62 mmol) and triphenylphosphine PPh_3 (1.44 g, 5.5 mmol) in THF (10 mL) at -15°C was added diethyl azodicarboxylate (DEAD) (1.11 g, 5.5 mmol), and this was stirred for 5 min. Diphenyl phosphorazidate (DPPA) (1.51 g, 5.5 mmol) was added, and the reaction mixture was stirred at 25 °C overnight. The solvent was removed, and the crude residue was purified by dry flash chromatography (pentane/ CH_2Cl_2 , 6:1) followed by flash chromatography (pentane/ CH_2Cl_2 , 10:1–1:1) to give the product as a white solid (0.72 g, 65%): mp 75–77 °C; IR (CH_2Cl_2) ν 3042, 2936, 2113, 1595, 1492 cm^{-1} ; $[\alpha]_{\text{D}} = -39.0^\circ$ ($c = 1.09$, CHCl_3 , 25 °C); $^1\text{H NMR}$ (270.2 MHz, CDCl_3) δ 7.29 (dd, $J = 8.6, 7.6$ Hz, 4H, OArH[m]), 6.98 (t, $J = 7.4$ Hz, 2H, OArH[p]), 6.92 (d, $J = 8.6$ Hz, 4H, OArH[o]), 4.33 (m, 2H, CHOC), 4.25 (d, $J = 6.3$ Hz, 4H, CH_2OPh), 3.75 (m, 2H, CHN_3), 1.46 (s, 6H, CH_3); $^{13}\text{C NMR}$ (67.8 MHz, CDCl_3) δ 157.9, 129.6, 121.6, 114.6, 110.9, 76.9, 67.8, 59.5, 26.9. Anal. ($\text{C}_{21}\text{H}_{24}\text{N}_6\text{O}_4$) C, H, N.

2,5-Diazido-1,6-di-O-phenyl-2,5-dideoxy-D-iditol (11a). **Method III.** Compound **10a** (2.0 g, 4.71 mmol) was added to a mixture of acetonitrile (60 mL) and 3 M HCl (15 mL). The temperature was raised to 50 °C and maintained for 4 h. The solvent was removed, and the crude product was purified by flash chromatography (CH_2Cl_2 to $\text{CH}_2\text{Cl}_2/\text{CH}_3\text{OH}$, 25:1) to give the product as a white solid (1.45 g, 80%); IR (film) ν 3427, 3064, 2932, 2110, 1595, 1493, 1239 cm^{-1} ; $[\alpha]_{\text{D}} = -25.1^\circ$ ($c = 1.0$, CHCl_3 , 25 °C); $^1\text{H NMR}$ (270.2 MHz, CDCl_3) δ 7.29 (dd, $J = 7.9, 6.6$ Hz, 4H, OArH[m]), 7.0 (t, $J = 7.3$ Hz, 2H, OArH[p]), 6.92 (d, $J = 7.9$ Hz, 4H, OArH[o]), 4.29 (m, 4H, CH_2OPh), 3.96 (m, 4H, CHOH , CHN_3), 2.91 (s, 2H, OH); $^{13}\text{C NMR}$ (67.8 MHz, CDCl_3) δ 157.9, 129.6, 121.7, 114.6, 70.9, 68, 62.4; HRMS calcd for $\text{C}_{18}\text{H}_{20}\text{N}_6\text{O}_4$ 384.1546, found 384.1544.

2,5-Diazido-3,4-bis-O-[(2-methoxyethoxy)methyl]-1,6-di-O-phenyl-2,5-dideoxy-D-iditol (12a). **Method IV.** Compound **11a** (1.39 g, 3.63 mmol) in THF (75 mL) was stirred with NaH (0.43 g, 14.5 mmol) for 15 min. (2-Methoxyethoxy)methyl chloride (MEMCl) (1.81 g, 14.5 mmol) was added, and the reaction mixture was stirred overnight. The reaction was quenched with aqueous saturated NH_4Cl ; the mixture was diluted with water and extracted with ether (2×100 mL). The combined ether extracts were dried and concentrated. The crude product was purified by flash chromatography (pentane/ CH_2Cl_2 , 1:1, to CH_2Cl_2) to give the product as a colorless oil (1.73 g, 85%): IR (film) ν 3041, 2927, 2104, 1599, 1498 cm^{-1} ; $[\alpha]_{\text{D}} = -29.6^\circ$ ($c = 1.0$, CHCl_3 , 25 °C); $^1\text{H NMR}$ (270.2 MHz, CDCl_3) δ 7.29 (dd, $J = 8.2, 7.6$ Hz, 4H, OArH[m]), 6.98 (t, $J = 7.2$ Hz, 2H, OArH[p]), 6.94 (d, $J = 8.2$ Hz, 4H, OArH[o]), 4.9 (d, $J = 6.9$ Hz, 2H, OCH_2O), 4.81 (d, $J = 6.9$ Hz, 2H, OCH_2O), 4.31 (m, 4H, CH_2OPh), 4.08 (m, 4H, CHOCH_2 , CHN_3), 3.78 (dt, $J = 11.2, 4.3$ Hz, 2H, $\text{OCH}_2\text{CH}_2\text{O}$), 3.69 (dt, $J = 11.6, 4.6$ Hz, 2H, $\text{OCH}_2\text{CH}_2\text{O}$), 3.51 (t, $J = 4.5$ Hz, 4H, $\text{OCH}_2\text{CH}_2\text{O}$), 3.35 (s, 6H, CH_3); $^{13}\text{C NMR}$ (67.8 MHz, CDCl_3) δ 158.1, 129.5, 121.4, 114.6, 97.3, 77.6, 71.6, 68.3, 68, 60.6, 59. Anal. ($\text{C}_{26}\text{H}_{36}\text{N}_6\text{O}_8$) C, H, N.

(4R,5S,6S,7R)-5,6-Bis[(2-methoxyethoxy)methoxy]-4,7-bis(phenoxymethyl)-1,3-diazepan-2-one (13a). **Method V.** To compound **12a** (716 mg, 1.28 mmol) in ethyl acetate (25 mL) was added a catalytic amount of 10% Pd/C. Hydrogen was added to the system at atmospheric pressure, and the reaction mixture was stirred overnight. The suspension was filtered through Celite and concentrated to give a colorless oil. In order to ensure high dilution conditions, the crude amine (595 mg) was dissolved in CH_2Cl_2 (50 mL) and the carbonyl-diimidazole (CDI) (230 mg, 1.41 mmol) was dissolved in CH_2Cl_2 (50 mL), respectively; this was slowly added with a syringe pump to a reservoir of CH_2Cl_2 (300 mL), and the reaction mixture was stirred overnight. The solvent was removed, and purification by flash chromatography (CH_2Cl_2 to $\text{CH}_2\text{Cl}_2/\text{CH}_3\text{OH}$, 100:1) gave the product as a white solid (426 mg, 67%): mp 218–220 °C; IR (KBr) ν 3228, 3106, 2936, 1687, 1600, 1498, 1466, 1243 cm^{-1} ; $[\alpha]_{\text{D}} = +50.1^\circ$ ($c = 1.14$, CHCl_3 , 25 °C); $^1\text{H NMR}$ (399.8 MHz, CDCl_3) δ 7.28 (dd, $J = 8.8, 7.5$ Hz, 4H, OArH[m]), 6.99 (t, $J = 7.4$ Hz, 2H, OArH[p]), 6.95 (d, $J = 8.8$ Hz, 4H, OArH[o]), 4.86 (d, $J = 7.1$ Hz, 2H, OCH_2O), 4.79 (d, $J = 7.3$ Hz, 2H, OCH_2O), 4.73 (br s, 2H, NH), 4.11 (app s, 6H, CHOCH_2 , CH_2OPh), 3.97 (br s, 2H, CHN), 3.75 (dt, $J = 11.3, 4.2$ Hz, 2H, $\text{OCH}_2\text{CH}_2\text{O}$), 3.71 (dt, $J = 11.5, 4.2$ Hz, 2H, $\text{OCH}_2\text{CH}_2\text{O}$), 3.52 (t, $J = 4.5$ Hz, 4H, $\text{OCH}_2\text{CH}_2\text{O}$), 3.35 (s, 6H, CH_3); $^{13}\text{C NMR}$ (67.8 MHz, CDCl_3) δ 164.0, 157.9, 129.5, 121.4, 114.6, 96.3, 74.6, 72.0, 67.8, 67.3, 58.9, 50.5. Anal. ($\text{C}_{27}\text{H}_{38}\text{N}_2\text{O}_9$) C, H, N.

(4R,5S,6S,7R)-1,3-Dibenzyl-5,6-bis[(2-methoxyethoxy)methoxy]-4,7-bis(phenoxymethyl)-1,3-diazepan-2-one (14a). **Method VI.** To compound **13a** (150 mg, 0.28 mmol) in DMF (10 mL) were added NaH (80% suspension) (50 mg, 1.68 mmol) and benzyl bromide (287 mg, 1.68 mmol). The reaction mixture was stirred under a N_2 atmosphere overnight. Excess NaH was quenched carefully with ethanol, and the reaction mixture was diluted with water and extracted with ether (2×50 mL). The combined ether extracts were washed with brine, dried, and concentrated. Purification by flash chromatography ($\text{CH}_2\text{Cl}_2/\text{CH}_3\text{OH}$, 100:1) gave a colorless oil (180 mg, 89%): IR (film) ν 3062, 2888, 1653, 1599, 1496, 1471 cm^{-1} ; $[\alpha]_{\text{D}} = -56.2^\circ$ ($c = 1.26$, CHCl_3 , 25 °C); $^1\text{H NMR}$ (270.2 MHz, CDCl_3) δ 7.25 (m, 14H, ArH), 6.95 (t, $J = 7.2$ Hz, 2H, OArH[p]), 6.82 (d, $J = 7.9$ Hz, 4H, OArH[o]), 5.16 (d, $J = 14.2$ Hz, 2H, CH_2Ph), 4.58 (d, $J = 7.0$ Hz, 2H, OCH_2O), 4.49 (d, $J = 6.9$ Hz, 2H, OCH_2O), 4.32 (app t, $J = 9.8$ Hz, 2H, CH_2OPh), 4.22 (dd, $J = 9.9, 2.6$ Hz, 2H, CH_2OPh), 4.09 (d, $J = 14.1$ Hz, 2H, CH_2Ph), 3.89 (br d, $J = 9.6, 2\text{H}$, CHN), 3.45–3.3 (m, 10H, CHOCH_2 , $\text{OCH}_2\text{CH}_2\text{O}$), 3.28 (s, 6H, CH_3); $^{13}\text{C NMR}$ (67.8 MHz, CDCl_3) δ 161.8, 158.6, 138.5, 129.6 (2 C), 128.9, 127.7, 121.2, 114.9, 96.5, 77.0, 71.7, 67.8, 65.7, 59.9, 59.1, 55.9. Anal. ($\text{C}_{41}\text{H}_{50}\text{N}_2\text{O}_9$) C, H, N.

(4R,5S,6S,7R)-1,3-Dibenzyl-4,7-bis(phenoxymethyl)-5,6-dihydroxy-1,3-diazepan-2-one (1a). **Method VII.** To a solution of acetonitrile (4 mL) and 3 M HCl (1 mL) was added **14a** (100 mg, 0.14 mmol). The reaction mixture was heated

to 50 °C for 5 h. The solvent was removed, and purification by flash chromatography (CH₂Cl₂ to CH₂Cl₂/CH₃OH, 100:1) gave a white solid (67.2 mg, 89%): mp 218–221 °C; IR (KBr) ν 3600–3300, 3026, 2892, 1586, 1496 cm⁻¹; [α]_D = -29.5° (*c* = 0.43, DMSO, 25 °C); ¹H NMR (270.2 MHz, DMSO-*d*₆) δ 7.31 (m, 14H, ArH), 6.96 (t, *J* = 7.3 Hz, 2H, OArH[*p*]), 6.86 (d, *J* = 8.1 Hz, 4H, OArH[*o*]), 5.36 (br s, 2H, OH), 4.96 (d, *J* = 14.3 Hz, 2H, CH₂Ph), 4.29 (m, 4H, CH₂OPh), 4.00 (d, *J* = 14.4 Hz, 2H, CH₂Ph), 3.70 (br m, 2H, CHN), 3.42 (m, 2H, CHOH); ¹³C NMR (67.8 MHz, DMSO-*d*₆) δ 161.8, 158.9, 139.4, 130.2, 129.5, 129.2, 127.9, 121.5, 115.3, 70.8, 66.0, 62.2, 56.2. Anal. (C₃₃H₃₄N₂O₅) C, H, N.

(4S,5R,6R,7S)-1,3-Dibenzyl-4,7-bis(phenoxyethyl)-5,6-dihydroxy-1,3-diazepan-2-one (1b). Compound **1b** was synthesized from **14b** according to method VII in 72% yield: mp 219–223 °C; IR (CHCl₃) ν 3585, 3056, 2937, 1645, 1596, 1463, 1449, 1269, 1237 cm⁻¹; [α]_D = +23.3° (*c* = 0.30, DMSO, 25 °C); ¹H NMR (270.2 MHz, DMSO-*d*₆) δ 7.3 (m, 14H, ArH), 6.95 (t, *J* = 7.3 Hz, 2H, OArH[*p*]), 6.85 (d, *J* = 8.1 Hz, 4H, OArH[*o*]), 5.39 (br s, 2H, OH), 4.97 (d, *J* = 14.2 Hz, 2H, CH₂-Ph), 4.3 (m, 4H, CH₂OPh), 4.0 (d, *J* = 15.0 Hz, 2H, CH₂Ph), 3.75 (br m, 2H, CHN), 3.37 (m, 2H, CHOH); ¹³C NMR (67.5 MHz, DMSO-*d*₆) δ 160.9, 158.0, 138.5, 129.4, 128.7, 128.4, 127.1, 120.7, 114.4, 69.8, 65.0, 61.3, 55.3. Anal. (C₃₃H₃₄N₂O₅) C, H, N.

(4S,5S,6S,7S)-1,3-Dibenzyl-4,7-bis(phenoxyethyl)-5,6-dihydroxy-1,3-diazepan-2-one (1c). Compound **1c** was synthesized from **14c** according to method VII in 76% yield: IR (KBr) ν 3600–3200, 3030, 2927, 1621, 1598, 1495, 1470, 1241 cm⁻¹; [α]_D = +47.3° (*c* = 0.69, CHCl₃, 25 °C); ¹H NMR (399.8 MHz, CDCl₃) δ 7.35 (m, 4H, ArH), 7.28 (m, 10H, ArH), 7.00 (t, *J* = 7.3 Hz, 2H, OArH[*p*]), 6.85 (d, *J* = 8.8 Hz, 4H, OArH[*o*]), 4.98 (d, *J* = 14.4 Hz, 2H, CH₂Ph), 4.13 (dd, *J* = 9.7, 5.3 Hz, 2H, CH₂OPh), 4.08 (dd, *J* = 9.7, 5.3 Hz, 2H, CH₂OPh), 4.02 (d, *J* = 14.4 Hz, 2H, CH₂Ph), 3.93 (m, 2H, CHOH), 3.42 (m, 2H, CHN), 2.62 (br s, 2H, OH); ¹³C NMR (67.5 MHz, CDCl₃) δ 163.3, 158.2, 137.8, 129.5, 129.2, 128.7, 127.7, 121.4, 114.6, 71.0, 66.9, 63.2, 52.9. Anal. (C₃₃H₃₄N₂O₅) C, H, N.

(4R,5R,6R,7R)-1,3-Dibenzyl-4,7-bis(phenoxyethyl)-5,6-dihydroxy-1,3-diazepan-2-one (1d). Compound **1d** was synthesized from **14d** according to method VII in 70% yield: IR (KBr) ν 3600–3200, 3030, 2926, 1621, 1598, 1494, 1469 cm⁻¹; [α]_D = -43.3° (*c* = 0.51, CHCl₃, 25 °C); ¹H NMR (399.8 MHz, CDCl₃) δ 7.37–7.25 (m, 14H, ArH), 7.00 (t, *J* = 7.3 Hz, 2H, OArH[*p*]), 6.85 (d, *J* = 7.8 Hz, 4H, OArH[*o*]), 4.99 (d, *J* = 14.4 Hz, 2H, CH₂Ph), 4.12 (dd, *J* = 9.6, 5.1 Hz, 2H, CH₂OPh), 4.09 (dd, *J* = 9.6, 5.1 Hz, 2H, CH₂OPh), 4.03 (d, *J* = 14.2 Hz, 2H, CH₂Ph), 3.95 (m, 2H, CHOH), 3.42 (m, 2H, CHN), 2.55 (br s, 2H, OH); ¹³C NMR (67.5 MHz, CDCl₃) δ 163.2, 158.2, 137.8, 129.5, 129.3, 128.7, 127.7, 121.4, 114.6, 71.0, 66.9, 63.2, 52.9. Anal. (C₃₃H₃₄N₂O₅) C, H, N.

1,6-Dibenzyl-3,4-O-isopropylidene-1,6-dideoxy-L-mannitol (15). Method VIII. To a stirred suspension of CuI (6.1 g, 32 mmol) in THF (50 mL) at -20 °C was added benzylmagnesium chloride (32 mL of a 2.0 M solution, 64 mmol). After 1 h at 0 °C, the diepoxide **8a** (2.0 g, 10.75 mmol) was added and the reaction mixture was stirred for 30 min at 0 °C. The reaction was quenched with saturated aqueous NH₄Cl; the mixture was diluted with water (50 mL) and extracted with ether (3 × 150 mL). The combined ether extracts were washed with brine (50 mL), dried, and concentrated. Purification by flash chromatography (CH₂Cl₂ to CH₂Cl₂/CH₃OH, 100:1) gave a white solid (3.47 g, 87%): mp 80–83 °C; IR (KBr) ν 3600–3100, 2922, 1600, 1494, 1453, 1374, 1069 cm⁻¹; [α]_D = -34.6° (*c* = 0.786, CHCl₃, 25 °C); ¹H NMR (270.2 MHz, CDCl₃) δ 7.2 (m, 10H, ArH), 3.67 (m, 4H, CHOH, CHOC), 3.24 (s, 2H, OH), 2.87 (ddd, *J* = 13.5, 10.2, 5.6 Hz, 2H, CH₂Ph), 2.74 (ddd, *J* = 13.9, 9.6, 6.6 Hz, 2H, CH₂Ph), 2.11 (dddd, *J* = 13.9, 10.2, 5.9, 4.0 Hz, 2H, CH₂CH₂Ph), 1.78 (dddd, *J* = 13.9, 9.3, 5.3, 5.3 Hz, 2H, CH₂CH₂Ph), 1.35 (s, 6H, CH₃); ¹³C NMR (67.8 MHz, CDCl₃) δ 141.9, 128.5, 128.4, 125.8, 108.9, 82.9, 72.5, 35.7, 31.3, 26.8. Anal. (C₂₃H₃₀O₄) C, H.

2,5-Diazido-1,6-dibenzyl-3,4-O-isopropylidene-1,2,5,6-tetradecyloxy-D-Iditol (16). Compound **16** was synthesized from **15** according to method II in 70% yield: IR (CH₂Cl₂) ν 3073, 2937, 2107, 1603, 1496, 1234 cm⁻¹; [α]_D = -47.7° (*c* =

1.42, CHCl₃, 25 °C); ¹H NMR (399.8 MHz, CDCl₃) δ 7.34–7.18 (m, 10H, ArH), 4.08 (app s, 2H, CHOC), 2.93 (dd, *J* = 9.8, 4.4 Hz, 2H, CHN₃), 2.85 (ddd, *J* = 13.9, 9.1, 5.4 Hz, 2H, CH₂Ph), 2.73 (ddd, *J* = 13.9, 8.8, 7.3 Hz, 2H, CH₂Ph), 2.14 (dddd, *J* = 14.4, 9.0, 9.0, 5.4 Hz, 2H, CH₂CH₂Ph), 1.93 (dddd, *J* = 14.2, 9.3, 7.3, 4.6 Hz, 2H, CH₂CH₂Ph), 1.48 (s, 6H, CH₃); ¹³C NMR (67.8 MHz, CDCl₃) δ 140.4, 128.6, 128.4, 126.2, 110.4, 79.7, 59.3, 32.4, 26.8. Anal. (C₂₃H₂₈N₆O₂) C, H, N.

2,5-Diazido-1,6-dibenzyl-1,2,5,6-tetradecyloxy-D-Iditol (17). Compound **17** was synthesized from **16** according to method III in 86% yield: mp 72–73 °C; IR (KBr) ν 3537, 3500–3200, 3030, 2941, 2103, 1602, 1495, 1454, 1280 cm⁻¹; [α]_D = +6.3° (*c* = 1.09, CHCl₃, 25 °C); ¹H NMR (270.2 MHz, CDCl₃) δ 7.33–7.25 (m, 4H, ArH), 7.23–7.15 (m, 6H, ArH), 3.61 (dt, *J* = 7.2, 4.0 Hz, 2H, CHOH), 3.29 (ddd, *J* = 9.5, 5.6, 3.9 Hz, 2H, CHN₃), 2.79 (ddd, *J* = 13.9, 11.2, 6.6 Hz, 2H, CH₂Ph), 2.72 (ddd, *J* = 13.9, 11.9, 7.6 Hz, 2H, CH₂Ph), 2.56 (d, *J* = 5.6 Hz, 2H, OH), 1.95 (m, 4H, CH₂CH₂Ph); ¹³C NMR (67.8 MHz, CDCl₃) δ 140.5, 128.6, 128.3, 126.2, 73.4, 62.9, 32.1, 32.0. Anal. (C₂₀H₂₄N₆O₂) C, H, N.

2,5-Diazido-1,6-dibenzyl-3,4-bis-O-[(2-methoxyethoxy)methyl]-1,2,5,6-tetradecyloxy-D-Iditol (18). Compound **18** was synthesized from **17** according to method IV in 70% yield: IR (film) ν 3026, 2926, 2103, 1603, 1496, 1453 cm⁻¹; [α]_D = -47.8° (*c* = 1.32, CHCl₃, 25 °C); ¹H NMR (399.8 MHz, CDCl₃) δ 7.3 (m, 4H, ArH), 7.21 (m, 6H, ArH), 4.90 (d, *J* = 7.1 Hz, 2H, OCH₂O), 4.81 (d, *J* = 6.8 Hz, 2H, OCH₂O), 3.82 (m, 2H, CHOCH₂), 3.76 (ddd, *J* = 11.0, 5.1, 3.6 Hz, 2H, OCH₂CH₂O), 3.64 (ddd, *J* = 10.7, 5.9, 3.7 Hz, 2H, OCH₂CH₂O), 3.46 (m, 4H, OCH₂CH₂O), 3.33 (s, 6H, CH₃), 3.18 (m, 2H, CHN₃), 2.80 (ddd, *J* = 13.9, 9.3, 5.6 Hz, 2H, CH₂Ph), 2.71 (ddd, *J* = 13.7, 9.3, 6.8 Hz, 2H, CH₂Ph), 2.1–1.93 (m, 4H, CH₂CH₂Ph); ¹³C NMR (67.8 MHz, CDCl₃) δ 140.7, 128.5, 128.4, 126.1, 97.3, 81.2, 71.5, 68.0, 60.7, 58.9, 32.4, 32.3. Anal. (C₂₈H₄₀N₆O₆) C, H, N.

(4R,5S,6S,7R)-5,6-Bis[(2-methoxyethoxy)methyl]-4,7-bis-(2-phenylethyl)-1,3-diazepan-2-one (19). Compound **19** was synthesized from **18** according to method V in 66% yield: mp 147–150 °C; IR (KBr) ν 3264, 3025, 2926, 1677, 1454, 1361 cm⁻¹; [α]_D = +30.9° (*c* = 1.25, CHCl₃, 25 °C); ¹H NMR (270.2 MHz, CDCl₃) δ 7.26 (m, 4H, ArH), 7.16 (m, 6H, ArH), 4.76 (d, *J* = 6.9 Hz, 2H, OCH₂O), 4.67 (d, *J* = 7.3 Hz, 2H, OCH₂O), 4.17 (br s, 2H, NH), 3.66 (br s, 2H, CHOCH₂), 3.53 (m, 6H, CHN, OCH₂CH₂O), 3.42 (m, 4H, OCH₂CH₂O), 3.31 (s, 6H, CH₃), 2.72 (m, 4H, CH₂Ph), 1.87 (m, 4H, CH₂CH₂Ph); ¹³C NMR (67.8 MHz, CDCl₃) δ 163.7, 140.6, 128.6, 128.4, 126.2, 96.3, 77.2, 71.5, 67.6, 59.0, 51.1, 33.9, 32.4. Anal. (C₂₉H₄₂N₂O₇) C, H, N.

(4R,5S,6S,7R)-1,3-Dibenzyl-5,6-bis[(2-methoxyethoxy)methyl]-4,7-bis-(2-phenylethyl)-1,3-diazepan-2-one (20). Compound **20** was synthesized from **19** according to method VI in 86% yield: IR (film) ν 3061, 2926, 1641, 1495, 1453, 1357 cm⁻¹; [α]_D = -60.9° (*c* = 1.17, CHCl₃, 25 °C); ¹H NMR (270.2 MHz, CDCl₃) δ 7.22 (m, 16H, ArH), 7.07 (m, 4H, ArH), 5.09 (d, *J* = 14.2 Hz, 2H, NCH₂Ph), 4.50 (d, *J* = 6.9 Hz, 2H, OCH₂O), 4.41 (d, *J* = 6.6 Hz, 2H, OCH₂O), 3.92 (d, *J* = 13.9 Hz, 2H, NCH₂Ph), 3.39 (m, 12H, CHOCH₂, CHN, OCH₂CH₂O), 3.32 (s, 6H, CH₃), 2.68 (ddd, *J* = 14.2, 9.2, 5.9 Hz, 2H, CH₂CH₂-Ph), 2.46 (dt, *J* = 14.2, 8.2 Hz, 2H, CH₂CH₂Ph), 1.96 (m, 4H, CH₂CH₂Ph); ¹³C NMR (67.8 MHz, CDCl₃) δ 126.3, 141.5, 138.1, 129.5, 128.6, 128.4, 128.3, 127.5, 125.9, 96.2, 77.1, 71.6, 67.1, 60.3, 59.0, 56.0, 33.0, 28.6. Anal. (C₄₃H₅₄N₂O₇) C, H, N.

(4R,5S,6S,7R)-1,3-Dibenzyl-4,7-bis-(2-phenylethyl)-5,6-dihydroxy-1,3-diazepan-2-one (2). Compound **2** was synthesized from **20** according to method VII in 91% yield: mp 210–212 °C; IR (KBr) ν 3600–3100, 3026, 2910, 1579, 1483, 1451, 1233 cm⁻¹; [α]_D = -22.6° (*c* = 1.17, CHCl₃, 23 °C); ¹H NMR (399.8 MHz, CDCl₃) δ 7.33–7.26 (m, 4H, ArH), 7.25–7.17 (m, 12H, ArH), 7.13 (m, 4H, ArH), 5.13 (d, *J* = 14.1 Hz, 2H, NCH₂Ph), 3.91 (d, *J* = 14.2 Hz, 2H, NCH₂Ph), 3.48 (s, 2H, CHOH), 3.35 (m, 2H, CHN), 2.78 (ddd, *J* = 13.2, 9.3, 6.1 Hz, 2H, CH₂CH₂Ph), 2.56 (ddd, *J* = 14.0, 9.8, 7.1 Hz, 2H, CH₂CH₂-Ph), 2.18 (s, 2H, OH), 2.10–1.96 (m, 4H, CH₂CH₂Ph); ¹³C NMR (67.8 MHz, CDCl₃) δ 162.4, 141.3, 137.9, 129.3, 128.5, 128.3, 128.2, 127.5, 125.9, 71.1, 60.7, 55.7, 33.1, 28.1. Anal. (C₃₅H₃₈N₂O₃) C, H, N.

1,6-Bis(2-phenylethyl)-3,4-O-isopropylidene-1,6-dideoxy- β -mannitol (21). Method IX. To Mg (763 mg, 31.8 mmol), ether (10 mL), and a small amount of iodine was added phenylethyl bromide (5.88 g, 31.8) under mild reflux. The phenylethylmagnesium bromide was added to a suspension of CuI (3.07 g, 16.1 mmol) in dry THF (35 mL) at -40°C . The temperature was slowly raised to -20°C over 60 min and then lowered to -40°C followed by addition of the diepoxide **8a** (1.0 g, 5.3 mmol). The temperature was once again slowly raised to -20°C . The reaction was quenched with saturated aqueous NH_4Cl ; the mixture was diluted with water (40 mL) and extracted with ether (3×150 mL). The combined ether extracts were washed with brine (40 mL), dried, and concentrated. Purification by flash chromatography (isohexane/ethyl acetate, 10:1–6:1) gave a white solid (1.37 g, 64%): IR (CHCl_3) ν 3600–3200, 3026, 2860, 1603, 1496, 1453 cm^{-1} ; $[\alpha]_{\text{D}} = -26.9^{\circ}$ ($c = 0.81$, CHCl_3 , 25°C); $^1\text{H NMR}$ (270.2 MHz, CDCl_3) δ 7.3–7.1 (m, 10H, ArH), 3.7–3.5 (m, 4H, CHOC, CHOH), 3.28 (br s, 2H, OH), 2.7–2.5 (m, 4H, $\text{CH}_2\text{CH}_2\text{Ph}$), 1.9–1.4 (m, 8H, $\text{CH}_2\text{CH}_2\text{CH}_2\text{Ph}$), 1.34 (s, 6H, CH_3); $^{13}\text{C NMR}$ (67.8 MHz, CDCl_3) δ 142.3, 128.3, 128.2, 125.7, 108.7, 83.0, 72.9, 35.8, 33.8, 26.8. Anal. ($\text{C}_{25}\text{H}_{34}\text{O}_4$) C, H.

2,5-Diazido-1,6-bis(2-phenylethyl)-3,4-O-isopropylidene-1,2,5,6-tetra-deoxy-D- β -iditol (22). Compound **22** was synthesized from **21** according to method II in 67% yield: IR (CHCl_3) ν 3050, 2974, 2111, 1602, 1496, 1453 cm^{-1} ; $[\alpha]_{\text{D}} = -46.2^{\circ}$ ($c = 1.05$, CHCl_3 , 25°C); $^1\text{H NMR}$ (270.2 MHz, CDCl_3) δ 7.28 (m, 4H, ArH), 7.18 (m, 6H, ArH), 3.99 (m, 2H, CHOC), 2.93 (m, 2H, CHN₃), 2.66 (t, $J = 7.2$ Hz, 4H, $\text{CH}_2\text{CH}_2\text{Ph}$), 1.93–1.55 (m, 8H, $\text{CH}_2\text{CH}_2\text{CH}_2\text{Ph}$), 1.44 (s, 6H, CH_3); $^{13}\text{C NMR}$ (67.8 MHz, CDCl_3) δ 141.5, 128.4, 128.3, 126.0, 110.3, 79.8, 60.5, 35.4, 30.2, 28.0, 26.8. Anal. ($\text{C}_{25}\text{H}_{32}\text{N}_6\text{O}_2$) C, H, N.

2,5-Diazido-1,6-bis(2-phenylethyl)-1,2,5,6-tetra-deoxy-D- β -iditol (23). Compound **23** was synthesized from **22** according to method III in 91% yield: mp 130 – 131°C ; $[\alpha]_{\text{D}} = +4.7^{\circ}$ ($c = 0.51$, DMSO, 25°C); IR (KBr) ν 3600–3200, 3026, 2936, 2110, 1495 cm^{-1} ; $^1\text{H NMR}$ (399.8 MHz, DMSO- d_6) δ 7.28 (m, 4H, ArH), 7.17 (m, 6H, ArH), 5.29 (d, $J = 7.8$ Hz, 2H, OH), 3.45 (m, 2H, CHOH), 3.32 (m, 2H, CHN₃), 2.58 (m, 4H, $\text{CH}_2\text{CH}_2\text{Ph}$), 1.74–1.42 (m, 8H, $\text{CH}_2\text{CH}_2\text{CH}_2\text{Ph}$); $^{13}\text{C NMR}$ (67.8 MHz, DMSO- d_6) δ 141.8, 128.2, 128.1, 125.6, 73.3, 62.5, 34.8, 29.2, 27.6. Anal. ($\text{C}_{22}\text{H}_{28}\text{N}_6\text{O}_2$) C, H, N.

2,5-Diazido-3,4-O-bis[(2-methoxyethoxy)methyl]-1,6-bis(2-phenylethyl)-1,2,5,6-tetra-deoxy-D- β -iditol (24). Compound **24** was synthesized from **23** according to method IV in 78% yield: IR (film) ν 3026, 2929, 2107, 1603, 1496, 1454 cm^{-1} ; $[\alpha]_{\text{D}} = -39.2^{\circ}$ ($c = 0.63$, CHCl_3 , 25°C); $^1\text{H NMR}$ (270.2 MHz, CDCl_3) δ 7.3–7.1 (m, 10H, ArH), 4.85 (d, $J = 6.9$ Hz, 2H, OCH_2O), 4.75 (d, $J = 6.9$ Hz, 2H, OCH_2O), 3.72 (m, 4H, $\text{OCH}_2\text{CH}_2\text{O}$), 3.61 (m, 2H, CHOCH_2), 3.48 (m, 4H, $\text{OCH}_2\text{CH}_2\text{O}$), 3.35 (s, 6H, CH_3), 3.17 (m, 2H, CHN₃), 2.65 (m, 4H, $\text{CH}_2\text{CH}_2\text{Ph}$), 1.72 (m, 8H, $\text{CH}_2\text{CH}_2\text{CH}_2\text{Ph}$); $^{13}\text{C NMR}$ (67.8 MHz, CDCl_3) δ 141.5, 128.3, 128.2, 125.8, 97.3, 81.1, 71.5, 68.0, 61.3, 58.9, 35.4, 30.0, 27.9. Anal. ($\text{C}_{30}\text{H}_{44}\text{N}_6\text{O}_6$) C, H, N.

(4R,5S,6S,7R)-5,6-Bis[(2-methoxyethoxy)methoxy]-4,7-bis(3-phenylpropyl)-1,3-diazepan-2-one (25). Compound **25** was synthesized from **24** according to method V in 78% yield: IR (CHCl_3) ν 3421, 3042, 2934, 1672, 1453 cm^{-1} ; $[\alpha]_{\text{D}} = +18.6^{\circ}$ ($c = 1.09$, CHCl_3 , 25°C); $^1\text{H NMR}$ (270.2 MHz, CDCl_3) δ 7.25 (m, 4H, ArH), 7.15 (m, 6H, ArH), 4.77 (d, $J = 7.6$ Hz, 2H, OCH_2O), 4.68 (d, $J = 7.6$ Hz, 2H, OCH_2O), 4.03 (br s, 2H, NH), 3.62 (m, 6H, CHOCH_2 , $\text{OCH}_2\text{CH}_2\text{O}$), 3.5 (br t, $J = 6.5$ Hz, 2H, CHNH), 3.43 (m, 4H, $\text{OCH}_2\text{CH}_2\text{O}$), 3.31 (s, 6H, CH_3), 2.63 (t, $J = 6.9$ Hz, 4H, $\text{CH}_2\text{CH}_2\text{Ph}$), 1.82–1.50 (m, 8H, $\text{CH}_2\text{CH}_2\text{CH}_2\text{Ph}$); $^{13}\text{C NMR}$ (67.8 MHz, CDCl_3) δ 164.0, 141.4, 128.4, 128.3, 126.0, 96.1, 76.6, 71.5, 67.7, 59.0, 51.5, 35.4, 32.0, 28.0. Anal. ($\text{C}_{31}\text{H}_{46}\text{N}_2\text{O}_7$) C, H, N.

(4R,5S,6S,7R)-1,3-Dibenzyl-5,6-bis[(2-methoxyethoxy)methoxy]-4,7-bis(3-phenylpropyl)-1,3-diazepan-2-one (26). Compound **26** was synthesized from **25** according to method VI in 78% yield: IR (film) ν 2927, 1641, 1495, 1453, 1110, 1035 cm^{-1} ; $[\alpha]_{\text{D}} = -55.7^{\circ}$ ($c = 0.67$, CHCl_3 , 25°C); $^1\text{H NMR}$ (270.2 MHz, CDCl_3) δ 7.25 (m, 14H, ArH), 7.12 (m, 6H, ArH), 4.96 (d, $J = 13.9$ Hz, 2H, NCH_2Ph), 4.50 (d, $J = 6.9$ Hz, 2H, OCH_2O), 4.42 (d, $J = 6.9$ Hz, 2H, OCH_2O), 3.83 (d, $J = 14.2$ Hz, 2H, NCH_2Ph), 3.37 (m, 12H, CHOCH_2 , CHN, $\text{OCH}_2\text{CH}_2\text{O}$),

2.42 (m, 4H, $\text{CH}_2\text{CH}_2\text{Ph}$), 1.80–1.33 (m, 8H, $\text{CH}_2\text{CH}_2\text{CH}_2\text{Ph}$); $^{13}\text{C NMR}$ (67.8 MHz, CDCl_3) δ 162.0, 141.9, 138.3, 129.6, 128.5, 128.3, 128.2, 127.4, 125.8, 96.3, 76.5, 71.6, 67.0, 61.3, 58.9, 56.0, 35.8, 28.9, 26.9. Anal. ($\text{C}_{45}\text{H}_{58}\text{N}_2\text{O}_7$) C, H, N.

(4R,5S,6S,7R)-1,3-Dibenzyl-4,7-bis(3-phenylpropyl)-5,6-dihydroxy-1,3-diazepan-2-one (3). Compound **3** was synthesized from **26** according to method VII in 74% yield: IR (film) ν 3600–3200, 3026, 2923, 1602, 1495, 1475, 1453 cm^{-1} ; $[\alpha]_{\text{D}} = -3.8^{\circ}$ ($c = 0.476$, CHCl_3 , 25°C); $^1\text{H NMR}$ (270.2 MHz, CDCl_3) δ 7.25 (m, 10H, ArH), 7.15 (m, 10H, ArH), 5.01 (d, $J = 14.2$ Hz, 2H, NCH_2Ph), 3.82 (d, $J = 14.2$ Hz, 2H, NCH_2Ph), 3.39 (br s, 2H, CHOH), 3.24 (br d, $J = 9.9$ Hz, 2H, CHN), 2.49 (m, 4H, $\text{CH}_2\text{CH}_2\text{Ph}$), 1.95 (br s, 2H, OH), 1.85–1.42 (m, 8H, $\text{CH}_2\text{CH}_2\text{CH}_2\text{Ph}$); $^{13}\text{C NMR}$ (67.8 MHz, CDCl_3) δ 162.2, 141.8, 138.1, 129.5, 128.6, 128.4, 128.3, 127.6, 125.9, 71.2, 61.8, 56.1, 35.9, 29.1, 26.5. Anal. ($\text{C}_{37}\text{H}_{42}\text{N}_2\text{O}_3$) C, H, N.

(4R,5S,6S,7R)-1,3-Bis{[4-methoxycarbonyl]phenyl}methyl}-5,6-bis[(2-methoxyethoxy)methoxy]-4,7-bis(phenoxymethyl)-1,3-diazepan-2-one (27). Compound **27** was synthesized from **13a** according to method VI in 72% yield, with methyl 4-(bromomethyl)benzoate as alkylating agent: IR (film) ν 3040, 2890, 1720, 1655, 1598, 1467 cm^{-1} ; $[\alpha]_{\text{D}} = -46.4^{\circ}$ ($c = 1.21$, CHCl_3 , 25°C); $^1\text{H NMR}$ (399.8 MHz, CDCl_3) δ 7.93 (d, $J = 8.3$ Hz, 4H, $\text{CH}_2\text{ArH}[\text{o}]$), 7.29 (m, 8H, ArH), 6.99 (t, $J = 7.3$ Hz, 2H, $\text{OArH}[\text{p}]$), 6.79 (d, $J = 8.0$ Hz, 4H, $\text{OArH}[\text{o}]$), 5.25 (d, $J = 14.2$ Hz, 2H, NCH_2Ar), 4.66 (d, $J = 6.9$ Hz, 2H, OCH_2O), 4.59 (d, $J = 7.1$ Hz, 2H, OCH_2O), 4.23 (d, $J = 6.3$ Hz, 4H, CH_2Ph), 4.18 (d, $J = 14.2$ Hz, 2H, NCH_2Ar), 3.93 (app s, 8H, CHN, CH_3OOC), 3.51–3.34 (m, 10H, CHOCH_2 , $\text{OCH}_2\text{CH}_2\text{O}$), 3.33 (s, 6H, CH_3); $^{13}\text{C NMR}$ (67.8 MHz, CDCl_3) δ 166.9, 161.4, 158.4, 143.7, 130.1, 129.6, 129.5, 129.4, 121.3, 114.8, 96.7, 71.7, 67.7, 65.5, 61.0, 59.1, 56.1, 52.2. Anal. ($\text{C}_{45}\text{H}_{54}\text{N}_2\text{O}_{13}$) C, H, N.

(4R,5S,6S,7R)-1,3-Bis{[4-(methoxycarbonyl)phenyl]methyl}-4,7-bis(phenoxymethyl)-5,6-dihydroxy-1,3-diazepan-2-one (6). Method X. To compound **27** (37 mg, 0.044 mmol) in methanol (4 mL) was added concentrated HCl (0.16 mL). The reaction mixture was heated to 40°C for 5 h. The solvent was removed. Purification by flash chromatography (CH_2Cl_2 to $\text{CH}_2\text{Cl}_2/\text{CH}_3\text{OH}$, 100:1) gave the product as a white solid (26.4 mg, 92%): mp 187 – 189°C ; IR (KBr) ν 3600–3300, 3040, 2950, 1720, 1604, 1471 cm^{-1} ; $[\alpha]_{\text{D}} = +3.2^{\circ}$ ($c = 1.43$, CHCl_3 , 25°C); $^1\text{H NMR}$ (399.8 MHz, CDCl_3) δ 7.95 (d, $J = 8.3$ Hz, 4H, $\text{CH}_2\text{ArH}[\text{o}]$), 7.31 (m, 8H, ArH), 7.02 (t, $J = 7.3$ Hz, 2H, $\text{OArH}[\text{p}]$), 6.8 (d, $J = 7.8$ Hz, 4H, $\text{OArH}[\text{o}]$), 5.15 (br d, $J = 11.9$ Hz, 2H, NCH_2Ar), 4.39 (m, 2H, CH_2OPh), 4.20 (m, 4H, NCH_2Ar , CH_2OPh), 3.92 (s, 6H, CH_3OOC), 3.80 (m, 2H, CHN), 3.59 (br s, 2H, CHOH), 2.8 (br s, 2H, OH); $^{13}\text{C NMR}$ (100.2 MHz, CDCl_3) δ 166.8, 161.5, 157.8, 143.3, 130.1, 129.7, 129.6, 129.2, 121.7, 114.6, 71.5, 65.1, 60.0, 55.8, 52.2. Anal. ($\text{C}_{37}\text{H}_{38}\text{N}_2\text{O}_9$) C, H, N.

(4R,5S,6S,7R)-1,3-Bis{[4-(hydroxymethyl)phenyl]methyl}-4,7-bis(phenoxymethyl)-5,6-dihydroxy-1,3-diazepan-2-one (7). Method XI. To compound **27** (60 mg, 0.072 mmol) in ether (2 mL) was added LiBH_4 (4.70 mg, 0.216 mmol). The reaction mixture was heated to reflux for 12 h. Excess LiBH_4 was quenched carefully with 1 M HCl (20 mL), and ether (20 mL) was added. The layers were separated, and the water phase was extracted with ether (2×20 mL). The combined ether extracts were washed with saturated NaHCO_3 (20 mL), dried, and concentrated to yield 45 mg of the crude product, which was used without further purification. To a solution of the crude diol in methanol (3 mL) was added concentrated HCl (1 mL); this was stirred overnight at 25°C . The solvent was removed, and the crude product was purified by flash chromatography ($\text{CH}_2\text{Cl}_2/\text{CH}_3\text{OH}$, 50:1–20:1) to give a white solid (27.1 mg, 63%): mp 161 – 163°C ; IR (KBr) ν 3600–3200, 3040, 2923, 1600, 1473, 1301 cm^{-1} ; $[\alpha]_{\text{D}} = -10.6^{\circ}$ ($c = 0.491$, CH_3OH , 25°C); $^1\text{H NMR}$ (399.8 MHz, CD_3OD) δ 7.34–7.18 (m, 12H, ArH), 6.98 (t, $J = 7.4$ Hz, 2H, $\text{OArH}[\text{p}]$), 6.90 (d, $J = 8.0$ Hz, 4H, $\text{OArH}[\text{o}]$), 5.12 (d, $J = 13.9$ Hz, 2H, NCH_2Ar), 4.58 (s, 4H, CH_2OH), 4.34 (m, 4H, CH_2OPh), 4.06 (d, $J = 14.2$ Hz, 2H, NCH_2Ar), 3.88 (m, 2H, CHN), 3.36 (br s, 2H, CHOH); $^{13}\text{C NMR}$ (100.2 MHz, CD_3OD) δ 163.4, 159.5, 141.8, 138.0, 130.1, 130.0, 128.0, 121.7, 115.4, 71.1, 65.8, 64.5, 62.3, 56.3. Anal. ($\text{C}_{35}\text{H}_{38}\text{N}_2\text{O}_7$) C, H, N.

(3R,4S,5S,6R)-4,5-Bis[(2-methoxyethyl)methoxy]-3,6-bis(phenoxyethyl)-1,2,7-thiadiazepane 1,1-Dioxide (28). Method XII. To compound **12a** (225 mg, 0.4 mmol) in ethyl acetate (5 mL) was added a catalytic amount of 10% Pd/C. Hydrogen was added to the system at atmospheric pressure, and the reaction mixture was stirred overnight. The suspension was filtered through Celite and concentrated to give a colorless oil. The crude amine (180 mg) was dissolved in dry pyridine (2.5 mL) under a N₂ atmosphere, and sulfamide (36 mg, 0.37 mmol) was added.²³ The reaction mixture was heated to reflux for 16 h. After cooling, the solution was diluted with ether (20 mL). The solution was washed with water (2 × 20 mL), then with 1 M aqueous HCl (2 × 10 mL), and finally with 10% aqueous Na₂CO₃ (10 mL). The ether solution was dried and concentrated. The residue was purified by flash chromatography (CH₂Cl₂ to CH₂Cl₂/CH₃OH, 19:1). This gave **29** as a pure white solid (138 mg, 60%): mp 72–74 °C; IR (mineral oil) ν 3356, 3278, 1587, 1490, 1350, 1251, 1226, 1162, 1125 cm⁻¹; [α]_D = -15.3° (c = 1.0, CHCl₃, 25 °C); ¹H NMR (270.2 MHz, CDCl₃) δ 7.29 (dd, *J* = 8.2, 7.9 Hz, 4H, OArH[m]), 6.98 (t, *J* = 7.3 Hz, 2H, OArH[p]), 6.88 (d, *J* = 8.3 Hz, 4H, OArH[o]), 5.35 (d, *J* = 10.9 Hz, 2H, NH), 4.83 (d, *J* = 7.3 Hz, 2H, OCH₂O), 4.78 (d, *J* = 7.3 Hz, 2H, OCH₂O), 4.3 (s, 2H, CHOCH₂), 4.13 (m, 2H, CH₂Oph), 4.04 (m, 2H, CH₂Oph), 3.90 (ddd, *J* = 8.6, 8.6, 5.5 Hz, 2H, CHN), 3.77 (m, 2H, OCH₂CH₂O), 3.61 (m, 2H, OCH₂CH₂O), 3.46 (m, 4H, OCH₂CH₂O), 3.32 (s, 6H, CH₃); ¹³C NMR (67.8 MHz, CDCl₃) δ 157.8, 129.6, 121.5, 114.5, 96.4, 74.1, 71.6, 68, 66.1, 59, 49.6. Anal. (C₂₆H₃₈N₂O₁₀S) C, H, N.

(3R,4S,5S,6R)-2,7-Dibenzyl-4,5-bis[(2-methoxyethoxy)methoxy]-3,6-bis(phenoxyethyl)-1,2,7-thiadiazepane 1,1-Dioxide (29). Method XIII. To compound **28** (30 mg, 0.053 mmol) in DMF (0.75 mL) was added NaH (80% suspension) (6.3 mg, 0.21 mmol). The reaction mixture was stirred at room temperature under a N₂ atmosphere for 30 min. Benzyl bromide (36 mg, 0.21 mmol) was added, and the reaction was allowed to proceed overnight. Excess NaH was quenched by dilution with water (5 mL), and the reaction mixture was extracted with ether (2 × 10 mL). The combined ether extracts were dried and concentrated. Purification by flash chromatography (CH₂Cl₂ to CH₂Cl₂/CH₃OH, 99:1) gave a colorless oil (39 mg, 99%): IR (film) ν 3052, 2927, 2886, 1600, 1497, 1329, 1243, 1157, 1112, 1036 cm⁻¹; [α]_D = -27.6° (c = 1.0, CHCl₃, 25 °C); ¹H NMR (270.2 MHz, CDCl₃) δ 7.48 (d, *J* = 6.9 Hz, 4H, ArH), 7.33 (dd, *J* = 7.6, 6.9 Hz, 4H, ArH), 7.22 (m, 6H, OArH[m], ArH), 6.93 (t, *J* = 7.3 Hz, 2H, OArH[p]), 6.68 (d, *J* = 8.6 Hz, 4H, OArH[o]), 5.03 (d, *J* = 17.2 Hz, 2H, NCH₂Ph), 4.84 (m, 6H, NCH₂Ph, OCH₂O), 4.47 (dd, *J* = 7.3, 6.6 Hz, 2H, CHN), 4.27 (s, 2H, CHOCH₂), 4.01 (m, 4H, CH₂Oph), 3.74 (m, 2H, OCH₂CH₂O), 3.67 (m, 2H, OCH₂CH₂O), 3.4 (m, 4H, OCH₂CH₂O), 3.28 (s, 6H, CH₃); ¹³C NMR (67.8 MHz, CDCl₃) δ 157.9, 139.6, 129.5, 128.6, 127.1, 127, 121.3, 114.4, 96, 76.8, 71.5, 67.9, 65.9, 59, 53.5, 52. Anal. (C₄₀H₅₀N₂O₁₀S) C, H, N.

(3R,4S,5S,6R)-2,7-Dibenzyl-3,6-bis(phenoxyethyl)-4,5-dihydroxy-1,2,7-thiadiazepane 1,1-Dioxide (4). Method XIV. To a solution of compound **29** (14 mg, 0.019 mmol) in methanol (1 mL) was added saturated HCl in ether (2 mL). The reaction mixture was stirred overnight at room temperature. The solvent was removed, and purification by flash chromatography (CH₂Cl₂ to CH₂Cl₂/CH₃OH, 49:1) gave a white solid (10 mg, 93%): mp 194–195 °C; IR (mineral oil) ν 3546, 3436, 1600, 1497, 1325, 1248, 1146 cm⁻¹; [α]_D = +7.4° (c = 1.0, CHCl₃, 25 °C); ¹H NMR (270.2 MHz, acetone-*d*₆) δ 7.54 (d, *J* = 7.2 Hz, 4H, ArH), 7.34–7.18 (m, 10H, ArH, OArH[m]), 6.91 (t, *J* = 7.3 Hz, 2H, OArH[p]), 6.79 (d, *J* = 7.9 Hz, 4H, OArH[o]), 5.05 (d, *J* = 17.5 Hz, 2H, NCH₂Ph), 4.96 (m, 4H, NCH₂Ph, OH), 4.45 (t, *J* = 6.6 Hz, 2H, CHN), 4.27 (m, 4H, CHOH, CH₂Oph), 4.16 (dd, *J* = 9.4, 6.3 Hz, 2H, CH₂Oph); ¹³C NMR (67.8 MHz, acetone-*d*₆) δ 159.8, 141.6, 130.8, 129.6, 128.5, 128.1, 122.3, 115.9, 75.3, 67.6, 56.4, 53.4. Anal. (C₃₂H₃₄N₂O₆S) C, H, N.

(3R,4S,5S,6R)-4,5-Bis[(2-methoxyethoxy)methoxy]-3,6-bis(2-phenylethyl)-1,2,7-thiadiazepane 1,1-Dioxide (30). Compound **30** was synthesized from **18** according to method XII in 77% yield: mp 98–99 °C; IR (KBr) ν 3208, 3029, 2917, 1604, 1495, 1456, 1418 cm⁻¹; [α]_D = +55.3° (c = 0.64, CHCl₃, 23 °C); ¹H NMR (399.8 MHz, CDCl₃) δ 7.30–7.16 (m, 10H,

ArH), 5.08 (d, *J* = 11.2 Hz, 2H, NH), 4.75 (d, *J* = 7.4 Hz, 2H, OCH₂O), 4.70 (d, *J* = 7.0 Hz, 2H, OCH₂O), 3.72 (app s, 2H, CHOCH₂), 3.6 (ddd, *J* = 11.0, 11.0, 3.9 Hz, 2H, CHN), 3.48 (m, 4H, OCH₂CH₂O), 3.41 (m, 4H, OCH₂CH₂O), 3.33 (s, 6H, CH₃), 2.88 (ddd, *J* = 10.2, 8.6, 5.2 Hz, 2H, CH₂CH₂Ph), 2.72 (ddd, *J* = 14.0, 8.0, 8.0 Hz, 2H, CH₂CH₂Ph), 1.89 (dddd, *J* = 13.9, 11.0, 8.8, 5.2 Hz, 2H, CH₂CH₂Ph), 1.66 (dddd, *J* = 12.2, 8.3, 8.3, 3.9 Hz, 2H, CH₂CH₂Ph); ¹³C NMR (100.2 MHz, CDCl₃) δ 141.6, 128.8, 128.5, 126.0, 96.8, 80.4, 71.6, 67.8, 59.0, 50.2, 35.4, 32.0. Anal. (C₂₈H₄₂N₂O₈S) C, H, N.

(3R,4S,5S,6R)-2,7-Dibenzyl-4,5-bis[(2-methoxyethoxy)methoxy]-3,6-bis(2-phenylethyl)-1,2,7-thiadiazepane 1,1-Dioxide (31). Compound **31** was synthesized from **30** according to method XIII in 88% yield: IR (KBr) ν 3062, 2924, 1603, 1496, 1452, 1308 cm⁻¹; [α]_D = -3.5° (c = 1.27, CHCl₃, 23 °C); ¹H NMR (399.8 MHz, DMSO-*d*₆) δ 7.45 (d, *J* = 7.1 Hz, 4H, ArH), 7.36 (app t, *J* = 7.8, 7.1 Hz, 4H, ArH), 7.30 (t, *J* = 7.3 Hz, 2H, ArH), 7.18 (app t, *J* = 7.5, 6.9 Hz, 4H, ArH), 7.11 (t, *J* = 7.3 Hz, 2H, ArH), 6.90 (d, *J* = 7.0 Hz, 4H, ArH), 4.82 (d, *J* = 16.6 Hz, 2H, NCH₂Ph), 4.73 (d, *J* = 6.9 Hz, 2H, OCH₂O), 4.69 (d, *J* = 6.6 Hz, 2H, OCH₂O), 5.53 (br m, 2H, NCH₂Ph), 3.81 (br s, 2H, CHOCH₂), 3.70 (br s, 2H, CHN), 3.52 (br m, 4H, OCH₂CH₂O), 3.39 (t, *J* = 4.6 Hz, 4H, OCH₂CH₂O), 3.19 (s, 6H, CH₃), 2.55 (br m, 2H, CH₂CH₂Ph), 2.33 (br m, 2H, CH₂CH₂Ph), 1.96 (br m, 2H, CH₂CH₂Ph), 1.85 (br m, 2H, CH₂CH₂Ph); ¹³C NMR (100.2 MHz, DMSO-*d*₆) δ 143.0, 140.8, 129.9, 129.8, 129.7, 129.4, 128.8, 127.4, 97.6, 81.1, 72.8, 68.8, 59.7, 58.2, 53.6, 34.2, 32.7. Anal. (C₄₂H₅₄N₂O₈S) C, H, N.

(3R,4S,5S,6R)-2,7-Dibenzyl-3,6-bis(2-phenylethyl)-4,5-dihydroxy-1,2,7-thiadiazepane 1,1-Dioxide (5). Compound **5** was synthesized from **31** according to method XIV in 77% yield: IR (KBr) ν 3600–3200, 3027, 2924, 1603, 1495, 1453, 1295 cm⁻¹; [α]_D = +47.9° (c = 1.21, CHCl₃, 23 °C); ¹H NMR (399.8 MHz, CDCl₃) δ 7.33 (d, *J* = 7.8 Hz, 4H, ArH), 7.23 (m, 6H, ArH), 7.11 (app t, *J* = 7.3 Hz, 4H, ArH), 7.06 (t, *J* = 7.3 Hz, 2H, ArH), 6.88 (d, *J* = 6.9 Hz, 4H, ArH), 4.68 (d, *J* = 15.6 Hz, 2H, NCH₂Ph), 4.28 (d, *J* = 15.6 Hz, 2H, NCH₂Ph), 3.70 (s, 2H, CHOH), 3.47 (d, *J* = 8.8 Hz, 2H, CHN), 2.68 (ddd, *J* = 13.2, 10.8, 4.9 Hz, 2H, CH₂CH₂Ph), 2.28 (br s, 2H, OH), 2.15 (ddd, *J* = 13.1, 9.8, 6.4 Hz, 2H, CH₂CH₂Ph), 1.98 (dddd, *J* = 14.6, 10.2, 10.2, 4.9 Hz, 2H, CH₂CH₂Ph), 1.85 (m, 2H, CH₂CH₂Ph); ¹³C NMR (67.8 MHz, CDCl₃) δ 141.8, 137.6, 128.85, 128.81, 128.7, 128.5, 128.1, 126.0, 74.1, 59.9, 53.9, 33.1, 29.4. Anal. (C₃₄H₃₈N₂O₄S) C, H, N.

Expression and Purification of HIV-1 Protease. HIV-1 protease was cloned and heterologously expressed in *Escherichia coli* and purified as described elsewhere.²⁸

Inhibition Studies. The degree of inhibition of HIV-1 protease was determined essentially as described previously²⁴ in a spectrophotometric assay using a chromophoric peptide, HIV substrate III (Bachem Feinchemicalien AG, Bubendorf, Switzerland). The measurements were performed in a total volume of 120 μ L by preincubating 60.8 μ M substrate and inhibitor in 50 mM glycine, 50 mM sodium acetate, 50 mM MES, 50 mM Tris-HCl, 1.0 M NaCl, 1 mM EDTA, and 1 mM DTT at pH 5.5 and 37 °C for 5 min. The reaction was started by addition of enzyme (30 nM), and the rate of cleavage was followed by continuously registering the change in absorbance at 300 nm. Inhibitors were dissolved in DMSO; the final concentration of DMSO was kept below 2.5%. Solubility was limited: high concentrations resulted in cloudiness in the assay cuvette and nonlinear traces of absorbance versus time. Inhibition was determined at 10 μ M for compounds not potent enough for IC₅₀-value determination. The IC₅₀-value is taken as the inhibitor concentration that results in 50% activity. It is estimated by plotting activity versus the logarithmic inhibitor concentration.

Determination of K_i-values was carried out by measuring the activity of 20 nM HIV-1 protease at different substrate and inhibitor concentrations. The mathematical analysis required an extensive data set comprising a wide range of substrate and inhibitor concentrations. The lowest substrate concentration that gave reproducible results and a detectable absorbance change was used (10 μ M). Due to limited solubility the highest concentration of substrate that could be used was 100 μ M. Two intermediate concentrations were also used (20

and 50 μM). The inhibitor concentration was chosen to be of the order of an initially estimated K_i -value and 2–4 times this concentration. At least three concentrations were used. Nanomolar concentrations of enzyme were used in order to obtain sufficiently rapid and accurate estimates of reaction velocities. Autoproteolysis of the enzyme and the sensitivity, which is limited by absorbance change upon cleavage of substrate, limit the time that can be used for the experiment, precluding the possibility of compensating lower rates achieved by decreasing the enzyme concentration with longer reactions. The total enzyme concentration is therefore of the same order as the concentration of inhibitor and requires the use of a rate equation that accounts for the presence of an enzyme–inhibitor complex for determination of K_i .²⁵ A rate equation for tight-binding inhibitors was fitted to the data by nonlinear regression analysis using QNFIT in SIMFIT.²⁹

Molecular Dynamics Simulations. MD simulations were performed with a modified version (Åqvist, unpublished) of the ENZYMIK program,³⁰ using the GROMOS force field²⁷ with modifications reported elsewhere.^{31,10} Two spherical simulation systems of radius 16 Å were set up for each inhibitor. In one, the inhibitor was placed in the center of the sphere, and SPC water molecules³² were added to fill the sphere. For the other simulation, the inhibitor was docked into the HIV proteinase, the simulation sphere was centered on the inhibitor, and water molecules were added to fill the nonprotein parts of the sphere. Protein atoms outside the simulation sphere were strongly restrained to their crystallographic positions and only interacted through bonds over the sphere boundary. The 16 Å radius sphere used in this work in fact contains most of the proteinase. MD simulations were then performed, first to allow the system to attain equilibrium and thereafter to collect data for the free energy calculations. For inhibitor–water simulations, the inhibitor was kept in the center of the sphere by restraining the carbonyl carbon. The simulation temperature was 300 K, the MD time step was 1 fs, and SCAAS surface restraints³³ were employed to hold the water surface at the desired radius. A nonbonded interaction cutoff radius of 10 Å was used for interactions between neutral noninhibitor groups, but the inhibitor and all charged groups interacted with everything inside the simulation sphere. Initial protein coordinates were from Brookhaven Protein Data bank entry 5HVP,^{5g} conformation 1. Net charges were left unmodified for the residues Asp 25, Arg 8, Arg 208, Asp 29, Asp 229, Asp 30, Asp 230, Arg 87, and Arg 287, while all other charged residues (far from the inhibitor) were replaced from the alternate set of GROMOS²⁷ residues with neutral dipolar groups. Asp 225 was considered to be protonated. Tetramethyl-substituted *RSSR* and *RRRR* cyclic urea rings were energy minimized by repeated simulated annealing using InsightII and Discover (Biosym/MSI, San Diego, CA), and the resulting minimum energy rings and their mirror images were used for modeling the ligands. The ligands were graphically docked into the protein, without manipulation of protein coordinates, to yield the starting models. We used Insight II (Biosym/MSI) for docking except for **3**, where the ligand was docked into the protein by a conformational search with the Macro Model program.³⁴

The systems consisting of the inhibitor in water were equilibrated for 190 ps at 300 K and then run for 125 ps, collecting data every 5 fs. The protein–inhibitor–water systems were heated under decreasing restraints in a 15 ps stepwise heating scheme, equilibrated for 175 ps at 300 K, and then run for 125 ps for data collection (250 ps for DMP 323 and **7**). Additional harmonic distance restraints of 4 kcal/mol Å² were applied during the heating and the first 25 ps of the equilibration to ensure correct initial geometries of the system. These were applied between the inhibitor carbonyl oxygen and the nitrogens of the peptide NH groups of Ile 50 and Ile 250 and between the inhibitor carbonyl carbon and the side chain carboxylate carbon of Asp 25 and Asp 225. For DMP 323 and **7**, similar additional restraints of 5 kcal/mol were applied between one carboxylate oxygen of aspartates 30 and 230 and the appropriate inhibitor hydroxyl group oxygen and hydrogen atoms, to ensure that the aspartates would find the possibility of hydrogen bonding to these hydroxyl groups. Otherwise, the

time needed for the aspartates to swing around to this equilibrium position from the crystallographic starting coordinates was very long. The data collected in inhibitor–water and inhibitor–protein–water simulations were the nonbonded force field interaction energies for inhibitor–surrounding interactions (where ‘surrounding’ refers to water or water + protein, respectively), as required by the LIE procedure¹⁰ used to evaluate the free energy (see below). Other details were as in the recently reported similar calculations on other HIV-1 proteinase inhibitors.¹¹

Free Energy Calculations. The LIE method is based on a linear response model of binding energetics. The simulation averages of the interaction energies between the inhibitor and its surroundings (no interactions within the inhibitor or within the surroundings are included in the average) are assumed to be linearly related to the free energy of solvation in that type of surrounding. The interaction energies are divided into electrostatic and van der Waals parts, with different proportionality constants for the different terms in the final expression for the binding energy below. As described elsewhere,^{10,35} these considerations result in the following semiempirical formula, where $\langle \rangle$ denotes MD time averages:

$$\Delta G_{\text{bind}} = \Delta G_{\text{el}} + \Delta G_{\text{vdw}} \approx \alpha (\langle V_{\text{vdw}} \rangle_{\text{prot}} - \langle V_{\text{vdw}} \rangle_{\text{wat}}) + \beta (\langle V_{\text{el}} \rangle_{\text{prot}} - \langle V_{\text{el}} \rangle_{\text{wat}}) \quad (1)$$

Here, α is an empirically determined constant, and β is predicted to be $1/2$ by the electrostatic linear response approximation.^{10,35} As in earlier work, we here use $\alpha = 0.161$ and $\beta = 1/2$ to estimate the binding free energies.

Acknowledgment. We gratefully acknowledge support from the Swedish Natural Science Research Council (NFR), Medivir AB, Huddinge, Sweden, and the Swedish National Board for Industrial and Technical Development (NUTEK). We wish to express our appreciation to Torleif Hård for valuable discussions.

Supporting Information Available: ¹H and ¹³C NMR spectral data, IR, optical rotations, and elemental analyses for compounds **9b–14b**, **9c–14c**, and **9d–14d** (6 pages). Ordering information is given on any current masthead page.

References

- (1) (a) Barré-Sinoussi, F.; Chermann, J. C.; Rey, F.; Nugeyre, M. T.; Chamaret, S.; Gruest, J.; Daugeat, C.; Axler-Blin, C.; Vézinet-Brun, F.; Rouzioux, C.; Rozenbaum, W.; Montagnier, L. Isolation of a T-lymphotropic retrovirus from a patient at risk for Acquired Immune Deficiency Syndrome (AIDS). *Science* **1983**, *220*, 868–871. (b) Gallo, R. C.; Sarin, P. S.; Gelman, E. P.; Robert-Guroff, M.; Richardson, E.; Kalyanaraman, V. S.; Mann, D.; Sidhu, G. D.; Stahl, R. E.; Zolla-Pazner, S.; Leibowitch, J.; Popovic, M. Isolation of Human T-Cell Leukemia Virus in Acquired Immune Deficiency Syndrome (AIDS). *Science* **1983**, *220*, 865–867. (c) Clavel, F.; Guétard, D.; Brun-Vézinet, F.; Chamaret, S.; Rey, M.-A.; Santos-Ferreira, M. O.; Laurent, A. G.; Daugeat, C.; Katlama, C.; Rouzioux, C.; Klatzmann, D.; Champalimaud, J. L.; Montagnier, L. Isolation of a new retrovirus from West African patients with AIDS. *Science* **1986**, *233*, 343–346.
- (2) Kohl, N. E.; Emini, E. A.; Schleif, W. A.; Davis, L. J.; Heimbach, J. C.; Dixon, R. A. F.; Scolnick, E. M.; Signal, I. S. Active Human Immunodeficiency Virus Protease is Required for Viral Infectivity. *Proc. Natl. Acad. Sci. U.S.A.* **1988**, *85*, 4686–4690.
- (3) (a) Tomasselli, A. G.; Howe, W. J.; Sawyer, T. K.; Wlodawer, A.; Heinrichson, R. L. The Complexities of AIDS: An Assessment of the HIV Protease as a Therapeutic Target. *Chim. Oggi* **1991**, *9*, 6–27. (b) Huff, J. R. HIV Protease: A Novel Chemotherapeutic Target for AIDS. *J. Med. Chem.* **1991**, *34*, 2305–2314. (c) Debouck, C. The HIV-1 Protease as a Therapeutic Target for AIDS. *AIDS Res. Human Retroviruses* **1992**, *8*, 153–164. (d) Darke, P. L.; Huff, J. R. HIV Protease as an Inhibitor Target for the Treatment of AIDS. In *Advances in Pharmacology*; August, J. T.; Anders, M. W.; Murad, F., Eds.; Academic Press: San Diego, CA, 1994; Vol. 25, pp 399–454.
- (4) (a) Martin, J. A. Recent Advances in the Design of HIV Protease Inhibitors. *Antiviral Res.* **1992**, *17*, 265–278. (b) Wlodawer, A.; Erickson, J. W. Structure-Based Inhibitors of HIV-1 Protease. *Annu. Rev. Biochem.* **1993**, *62*, 543–585. (c) Thairisrivongs, S. HIV Protease Inhibitors. In *Annual Reports in Medicinal Chemistry*; Bristol, J. A., Ed.; Academic Press: San Diego, CA,

- 1994; Vol. 29, pp 133–144. (d) De Clercq, E. Toward Improved Anti-HIV Chemotherapy: Therapeutic Strategies for Intervention with HIV Infections. *J. Med. Chem.* **1995**, *38*, 2491–2517. (e) Martin, J. A.; Redshaw, S.; Thomas, G. J. Inhibitors of HIV Proteinase. In *Progress in Medicinal Chemistry*; Ellis, G. P., Luscombe, D. K., Eds.; Elsevier Science: New York, 1995; Vol. 32, pp 239–287.
- (5) For early examples, see: (a) Wlodawer, A.; Miller, M.; Jaskólski, M.; Sathyanarayana, B. K.; Baldwin, E.; Weber, I. T.; Selk, L. M.; Clawson, L.; Schneider, J.; Kent, S. B. H. Conserved Folding in Retroviral Proteases: Crystal Structure of a Synthetic HIV-1 Protease. *Science* **1989**, *245*, 616–621. (b) Lapatto, R.; Blundell, T.; Hemmings, A.; Overington, J.; Wilderspin, A.; Wood, S.; Merson, J. R.; Whittle, P. J.; Danley, D. E.; Geoghegan, K. F.; Hawrylik, S. J.; Lee, S. E.; Scheld, K. G.; Hobart, P. M. X-ray analysis of HIV-1 proteinase at 2.7 Å resolution confirms structural homology among retroviral enzymes. *Nature* **1989**, *342*, 299–302. (c) Cooper, J. B.; Foundling, S. I.; Blundell, T. L.; Boger, J.; Jupp, R. A.; Kay, J. X-ray Studies of Aspartic Proteinase-Statine Inhibitor Complexes. *Biochemistry* **1989**, *28*, 8596–8603. (d) Miller, M.; Schneider, J.; Sathyanarayana, B. K.; Toth, M. V.; Marshall, G. R.; Clawson, L.; Selk, L.; Kent, S. B. H.; Wlodawer, A. Structure of Complex of Synthetic HIV-1 Protease with a Substrate-Based Inhibitor at 2.3 Å Resolution. *Science* **1989**, *246*, 1149–1152. (e) Navia, M. A.; Fitzgerald, P. M. D.; McKeever, B. M.; Leu, C.-T.; Heimbach, J. C.; Herber, W. K.; Sigal, I. S.; Darke, P. L.; Springer, J. P. Three-Dimensional Structure of Aspartyl Protease from human Immunodeficiency Virus HIV-1. *Nature* **1989**, *337*, 615–620. (f) 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.; Helfrich, R.; Paul, D. A.; Knigge, M. Design, Activity, and 2.8 Å Crystal Structure of a C₂-symmetric Inhibitor Complexed to HIV-1 Protease. *Science* **1990**, *249*, 527–533. (g) Fitzgerald, P. M. D.; McKeever, B. M.; VanMiddlesworth, J. F.; Springer, J. P.; Heimbach, J. C.; Leu, C.-T.; Herber, W. K.; Dixon, R. A. F.; Darke, P. L. Crystallographic Analysis of a Complex between Human Immunodeficiency Virus Type 1 Protease and Acetylpepstatin at 2.0-Å Resolution. *J. Biol. Chem.* **1990**, *265*, 14209–14219. (h) Swain, A. L.; Miller, M. M.; Green, J.; Rich, D. H.; Schneider, J.; Kent, S. B. H.; Wlodawer, A. X-ray Crystallographic Structure of a Complex Between a Synthetic Protease of Human Immunodeficiency Virus 1 and a Substrate-Based Hydroxyethylene Inhibitor. *Proc. Natl. Sci. U.S.A.* **1990**, *87*, 8805–8809. (i) Jaskólski, M.; Tomasselli, A. G.; Sawyer, T. K.; Staples, D. G.; Heinrichson, R. L.; Schneider, J.; Kent, S. B. H.; Wlodawer, A. Structure at 2.5-Å Resolution of Chemically Synthesized Human Immunodeficiency Virus Type 1 Protease Complexed with a Hydroxyethylene-Based Inhibitor. *Biochemistry* **1991**, *30*, 1600–1609. (j) Graves, B. J.; Hatada, M. H.; Miller, J. K.; Graves, M. C.; Roy, S.; Cook, C. H.; Kröhn, A.; Martin, J. A.; Roberts, N. A. The Three-Dimensional X-Ray Crystal Structure of HIV-1 Protease Complexed with a Hydroxyethylene Inhibitor. In *Structure and function of the Aspartic Proteinase*; Dunn, B. M., Ed.; Plenum Press: New York, 1991; pp 455–460. (k) Dreyer, G.; Lambert, D. M.; Meek, T. D.; Carr, T. J.; Tomaszek, T. A., Jr.; Fernandez, A. V.; Bartus, H.; Cacciavillani, E.; Hassell, A. M.; Minnich, M.; Petteway, S. R., Jr.; Metcalf, B. W. Hydroxyethylene Isostere Inhibitors of Human Immunodeficiency Virus-1 Protease: Structure-Activity Analysis Using Enzyme Kinetics, X-ray Crystallography, and Infected T-Cell Assays. *Biochemistry* **1992**, *31*, 6646–6659. (l) Tucker, T. J.; Lumma, W. C., Jr.; Payne, L. S.; Wai, J. M.; de Solmes, S. J.; Giuliani, E. A.; Darke, P. L.; Heimbach, J. C.; Zugay, J. A.; Schleich, W. A.; Quintero, J. C.; Emini, E. A.; Huff, J. R.; Anderson, P. S. A Series of Potent HIV-1 Protease Inhibitors Containing a Hydroxyethyl Secondary Amine Transition State Isostere: Synthesis, Enzyme Inhibition, and Antiviral Activity. *J. Med. Chem.* **1992**, *35*, 2525–2533. (m) Bone, R.; Vacca, J. P.; Anderson, P. S.; Holloway, M. K. X-ray Crystal Structure of the HIV Protease Complex with L-700,417, an Inhibitor with Pseudo C₂ symmetry. *J. Am. Chem. Soc.* **1992**, *113*, 9382–9384.
- (6) (a) Lam, P. Y. S.; Eyer mann, C. J.; Hodge, C. N.; Jadhav, P. K.; DeLucca, G. V. Patent WO 93/07128, 1993. (b) Lam, P. Y. S.; Jadhav, P. K.; Eyer mann, C. J.; Hodge, C. N.; DeLucca, G. V.; Rodgers, J. D. Patent WO 94/19329, 1994. (c) Lam, P. Y. S.; Jadhav, P. K.; Eyer mann, C. J.; Hodge, C. N.; Ru, Y.; Bachelier, L. T.; Meek, J. L.; Otto, M. J.; Rayner, M. M.; Wong, Y. N.; Chang, C.-H.; Weber, P. C.; Jackson, D. A.; Sharpe, T. R.; Erickson-Viitanen, S. Rational Design of Potent, Bioavailable, Nonpeptide Cyclic Ureas as HIV Protease Inhibitors. *Science* **1994**, *263*, 380–384. For related cyclic structures, see: (d) Chenera, B.; DesJarlais, R. L.; Finkelstein, J. A.; Eggleston, D. S.; Meek, T. D.; Tomaszek, T. A., Jr.; Dreyer, G. B. Nonpeptide HIV Protease Inhibitors Designed to Replace a Bound Water. *Bioorg. Med. Chem. Lett.* **1993**, *3*, 2717–2722. (e) Peyman, A.; Stahl, W.; Wagner, K.; Rubbert, D.; Budt, K.-H. Non-Peptide-Based Inhibitors of Human Immunodeficiency Virus-1 Protease. *Bioorg. Med. Chem. Lett.* **1994**, *4*, 2601–2604. (f) Randad, R. S.; Pan, W.; Gulnik, S. V.; Burt, S.; Erickson, J. W. De Novo Design of Nonpeptidic HIV-1 Protease Inhibitors: Incorporation of Structural Water. *Bioorg. Med. Chem. Lett.* **1994**, *4*, 1247–1252.
- (7) Sham, H. L.; Zhao, C.; Stewart, K. D.; Betebenner, D. A.; Lin, S.; Park, C. H.; Kong, X.-P.; Rosenbrook, W., Jr.; Herrin, T.; Madigan, D.; Vasavanonda, S.; Lyons, N.; Molla, A.; Saldívar, A.; Mersh, K. C.; McDonald, E.; Wideburg, N. E.; Denissen, J. F.; Robins, T.; Kempf, D. J.; Plattner, J. J.; Norbeck, D. W. A Novel Picomolar Inhibitor of Human Immunodeficiency Virus Type 1 Protease. *J. Med. Chem.* **1996**, *39*, 392–397.
- (8) (a) Tomioka, N.; Itai, A.; Iitaka, Y. A. Method for Fast Energy Estimation and Visualization of Protein-Ligand Interaction. *J. Comput.-Aided Mol. Des.* **1987**, *1*, 197–210. (b) Meng, E. C.; Shoichet, B. K.; Kuntz, I. D. Automated Docking with Grid-based Energy Evaluation. *J. Comput. Chem.* **1992**, *13*, 505–524. (c) Rotstein, S. H.; Murcko, M. A. GroupBuild: a Fragment-Based Method for De Novo Drug Design. *J. Med. Chem.* **1993**, *36*, 1700–1710. (d) Böhm, H.-J. The Development of a Simple Empirical Scoring Function to Estimate the Binding Constant for a Protein-Ligand Complex of Known Three-Dimensional Structure. *J. Comput.-Aided Mol. Des.* **1994**, *8*, 243–256.
- (9) Viswanadhan, V. N.; Reddy, M. R.; Wlodawer, A.; Varney, M. D.; Weinstein, J. N. An Approach to Rapid Estimation of Relative Binding Affinities of Enzyme Inhibitors: Application to Peptidomimetic Inhibitors of the Human Immunodeficiency Virus type 1 Protease. *J. Med. Chem.* **1996**, *39*, 705–712.
- (10) (a) Åqvist, J.; Medina, C.; Samuelsson, J.-E. A New Method for Predicting Binding Affinity in Computer-Aided Drug Design. *Protein Eng.* **1994**, *7*, 385–391. (b) Åqvist, J.; Mowbray, S. L. Sugar Recognition by a glucose/galactose receptor: evaluation of binding energetics from molecular dynamics simulations. *J. Biol. Chem.* **1995**, *270*, 9978–9981. (c) Åqvist, J. Calculation of absolute binding free energies for charged ligands and the effects of long-range electrostatic interactions. *J. Comput. Chem.* **1996**, *17*, 1587–1597.
- (11) Hansson, T.; Åqvist, J. Estimation of Binding Free Energies for HIV Proteinase Inhibitors by Molecular Dynamics Simulations. *Protein Eng.* **1995**, *8*, 1137–1144.
- (12) (a) Ghosh, A. K.; McKee, S. P.; Thompson, W. J. Stereocontrolled Synthesis of HIV-1 Protease Inhibitors with C₂-Axis of Symmetry. *Tetrahedron Lett.* **1991**, *32*, 5729–5732. (b) Chenera, B.; Boehm, J. C.; Dreyer, G. B. Synthesis of C₂-Symmetric and Pseudosymmetric HIV-1 Protease Inhibitors from D-Mannitol and D-Arabitol. *Bioorg. Med. Chem. Lett.* **1991**, *1*, 219–222. (c) Jadhav, P. K.; Woerner, F. J. Synthesis of C₂-Symmetric HIV-1 Protease Inhibitors from D-Mannitol. *Bioorg. Med. Chem. Lett.* **1992**, *2*, 353–356. (d) Dreyer, G. B.; Boehm, J. C.; Chenera, B.; DesJarlais, R. L.; Hassell, A. M.; Meek, T. D.; Tomaszek, T. A., Jr.; Lewis, M. A. Symmetric Inhibitor Binds HIV-1 Protease Asymmetrically. *Biochemistry* **1993**, *32*, 937–947. (e) Budt, K.-H.; Peyman, A.; Hansen, J.; Knolle, J.; Meichsner, C.; Paessens, A.; Ruppert, D.; Stowasser, B. HIV Protease Inhibitor HOE/BAY 793, Structure-Activity Relationships in a Series of C₂-Symmetric Diols. *Bioorg. Med. Chem.* **1995**, *3*, 559–571.
- (13) During the preparation of this manuscript the synthesis of a series of related compounds using a similar reaction sequence was reported: Nugiel, D. A.; Jacobs, K.; Worley, T.; Patel, M.; Kaltenbach, R. F., III; Meyer, D. T.; Jadhav, P. K.; De Lucca, G. V.; Smyser, T. E.; Klabe, R. M.; Bachelier, L. T.; Rayner, M. M.; Seitz, S. P. Preparation and Structure-Activity Relationship of Novel P1/P1'-Substituted Cyclic Urea-Based Human Immunodeficiency Virus Type-1 Protease Inhibitors. *J. Med. Chem.* **1996**, *39*, 2156–2169.
- (14) (a) Le Merrer, Y.; Duréault, A.; Greck, C.; Micas-Languin, D.; Gravier, C.; Depeyaz, J.-C. Synthesis of Diepoxides and Diaziridines, Precursors of Enantiomerically Pure α-Hydroxy and α-Amino Aldehydes or Acids, from D-Mannitol. *Heterocycles* **1987**, *25*, 541–548. (b) Wiggins, L. F. The Anhydrides of Polyhydric Alcohols. Part II. Derivatives of 1:2–5:6-Dianhydro Mannitol. *J. Chem. Soc.* **1946**, 384–388.
- (15) (a) Sexton, A. R.; Britton, E. C. Synthesis and Identification of Propylene Glycol Phenyl Ethers. *J. Am. Chem. Soc.* **1948**, *70*, 3606–3607. (b) Lipshutz, B. H.; Saumitra, S. Organocopper Reagents: Substitution, conjugate addition Carbo/Metallo-occupation, and other Reactions. In *Organic Reactions*; Paquette, L. A., Ed.; John Wiley & Sons: New York, 1992; Vol. 41, pp 135–631. (c) McSweeney, G. P.; Wiggins, L. F.; Wood, D. J. C. Anhydrides of Polyhydric Alcohols. Part XVI. The Action of Phenols on Some Ethylene Oxide Derivatives. *J. Chem. Soc.* **1952**, 37–53.
- (16) (a) Hughes, D. L. The Mitsunobu Reaction. In *Organic Reactions*; Paquette, L. A., Ed.; John Wiley & Sons: New York, 1992; Vol. 42, pp 335–656. (b) Mitsunobu, O. The Use of Diethyl Azodicarboxylate and Triphenylphosphine in Synthesis and Transformation of Natural Products. *Synthesis* **1981**, 1–28.

- (17) For a successful cyclization of a related *trans*-fused five-membered acetonide, see: Rossano, L. T.; Lo, Y. S.; Lee, Y.-C.; Meloni, D. J.; Moore, J. R.; Gale, T. M.; Arnett, J. F. A Practical Synthesis of Nonpeptide Cyclic Ureas as Potent HIV Protease Inhibitors. *Tetrahedron Lett.* **1995**, *36*, 4967–4970.
- (18) Corey, E. J.; Gras, J.-L.; Ulrich, P. A New General Method for Protection of the Hydroxyl Function. *Tetrahedron Lett.* **1976**, *809*–812.
- (19) (a) Sartori, E.; Camy, F.; Teulon, J. M.; Camborde, F.; Meignen, J.; Hertz, F.; Cloarec, A. Synthesis and analgesic activities of urea derivatives of α -amino-*N*-pyridyl benzene propanamide. *Eur. J. Med. Chem.* **1994**, *29*, 431–439. (b) Staab, H. A. Reaktionsfähige Heterocyclische Diamide der Kohlensäure. *Liebigs Ann. Chem.* **1957**, *609*, 75–83.
- (20) Wilkerson, W. W.; Hollins, A. Y.; Cheatham, W. W.; Lam, G. N.; Erickson-Viitanen, S.; Bachelier, L.; Cordova, B. C.; Klabe, R. M.; Meek, J. L. A Bis-[*N*-3-(1-Hydroxy-1-Methyl-Ethyl)-Benzyl]-Cyclic Urea as a HIV Protease Inhibitor. *Bioorg. Med. Chem. Lett.* **1995**, *5*, 3027–3032.
- (21) For a recent report on a stereoselective synthesis of DMP 323 from *D*-phenylalanine and a detailed discussion on bis-*N*-benzylation, see: Pierce, M. E.; Harris, G. D.; Islam, Q.; Radescia, L. A.; Storace, L.; Waltermir, R. E.; Wat, E.; Jadhav, P. K.; Emmet, G. C. Stereoselective Synthesis of HIV-1 Protease Inhibitor, DMP 323. *J. Org. Chem.* **1996**, *61*, 444–450.
- (22) Soai, K.; Ookawa, A. Mixed Solvents Containing Methanol as Useful Reaction Media for Unique Chemoselective Reductions with Lithium Borohydride. *J. Org. Chem.* **1986**, *51*, 4000–4005.
- (23) (a) Arya, V. P.; Shenoy, S. J. Synthesis of New Heterocycles: Part XV-Synthesis of Novel Cyclic & Acyclic Sulphamides. *Ind. J. Chem.* **1976**, *14B*, 766–769. (b) Jadhav, P. K.; Woerner, F. J. Synthesis of 8-Membered Cyclic Sulfamides: Novel HIV-1 Protease Inhibitors. *Tetrahedron Lett.* **1995**, *36*, 6383–6386.
- (24) Nillroth, U.; Besidsky, Y.; Classon, B.; Chattopadhyaya, J.; Ugi, I.; Danielson, U. H. Specific Interaction between HIV-1 proteinase and 5'-phosphate peptidomimetic derivatives of nucleoside analogues. *Drug. Des. Discovery* **1995**, *13*, 43–54.
- (25) Sculley, M. J.; Morrison, J. F. The Determination of Kinetic Constants Governing the Slow, Tight-Binding Inhibition of Enzyme-Catalysed Reactions. *Biochim. Biophys. Acta* **1986**, *874*, 44–53.
- (26) For a discussion on the protonation state of the catalytic aspartates, see: (a) Hyland, L. J.; Tomaszek, T. A., Jr.; Meek, T. D. Human Immunodeficiency Virus-1 Protease. 2. Use of pH Rate Studies and Solvent Kinetic Isotope Effects to Elucidate Details of Chemical Mechanism. *Biochemistry* **1992**, *30*, 8454–8463. (b) Davies, D. R. The Structure and Function of the Aspartic Proteinases. *Annu. Rev. Biophys. Chem.* **1990**, *19*, 189–215. (c) Yamazaki, T.; Nicholson, L. K.; Torchia, D. A.; Wingfield, P.; Stahl, S. J.; Kaufman, J. D.; Eyermann, C. J.; Hodge, C. N.; Lam, P. Y. S.; Ru, Y.; Jadhav, P. K.; Chang, C.-H.; Weber, P. C. NMR and X-ray evidence that the HIV protease catalytic aspartyl groups are protonated in the complex formed by the protease and a non-peptide cyclic urea-based inhibitor. *J. Am. Chem. Soc.* **1994**, *116*, 10791–10792.
- (27) van Gunsteren, W. F.; Berendsen, H. J. C. *Groningen Molecular simulation (GROMOS) Library Manual*; Biomos B.V.: Groningen, 1987.
- (28) Markgren, P.-O.; Nillroth, U.; Lindgren, M. T.; Danielson, U. H. Inhibition of HIV-1 Proteinase by Cu²⁺: Evidence for high affinity complex with immobilized Cu²⁺ and allosteric interactions. Submitted for publication.
- (29) Bardsley, W. G.; McGinlay, P. B.; Roig, M. G. Optimal Design for Model Discrimination Using F-test with Non Linear Biochemical Models. Criteria for Choosing the Number and Spacing of Experimental Points. *J. Theor. Biol.* **1989**, *139*, 85–102.
- (30) Warshel, A.; Creighton, S. Microscopic Free Energy Calculations in Solvated Macromolecules as a Primary Structure-Function Correlator and the MOLARIS Program. In *Computer simulation of biomolecular systems: theoretical and experimental approaches*; van Gunsteren, W. F.; Wiener, P. K., Eds.; ESCOM Science Publishers B.V.: Leiden, 1989; pp 120–138.
- (31) Åqvist, J.; Fothergill, M.; Warshel, A. Computer Simulation of the CO₂/HCO₃⁻ Interconversion Step in Human Carbonic Anhydrase I. *J. Am. Chem. Soc.* **1993**, *115*, 631–635.
- (32) Berendsen, H. J. C.; Postma, J. P. M.; van Gunsteren, W. F.; Hermans, J. Interaction Models for Water in Relation to Protein Hydration. In *Intermolecular Forces*; Pullman, B., Ed.; Reidel: Dordrecht, 1981; pp 331–342.
- (33) King, G.; Warshel, A. A Surface Constrained All-Atom Solvent Model for Effective Simulations of Polar Solutions. *J. Chem. Phys.* **1989**, *91*, 3647–3661.
- (34) Mohamadi, F.; Richards, N. G. J.; Guida, W. C.; Liskamp, R.; Lipton, M.; Caufield, C.; Chang, G.; Hendrickson, T.; Still, W. C. MacroModel-An Integrated Software System for Modeling Organic and Bioorganic Molecules Using Molecular Mechanics. *J. Comput. Chem.* **1990**, *11*, 440–467.
- (35) Åqvist, J.; Hansson, T. On the Validity of Electrostatic Linear Response in Polar Solvents. *J. Phys. Chem.* **1996**, *100*, 9512–9521.

JM960728J