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Chiral Recognition of Amino Acid Derivatives by 1,1'-Binaphthalene-8,8'-diol

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Abstract: Optically active 1,1'-binaphthalene-8,8'-diol is found to bind a variety of amines in CDCl₃ or C_6D_6 solution. Significant chiral recognition ($\Delta\Delta G^{\circ} = \sim 1.2$ kcal/mol) was observed in the valine derivative 4. A three-point binding motif is assumed. Copyright © 1996 Elsevier Science Ltd

Chiral recognition of organic substrates through non-covalent host-guest interaction has attracted increasing attention in recent years, since the process is directly applicable to chromatographic, 1 transporting, 2 catalytic, 3 and chemosensory 4 methodology. Several examples have been reported of chiral recognition of amino acids 5, sugars 4, ammonium ions 6, and organic acids 7 using well-designed organic host molecules. We have been interested in developing the ability of 1,1'-binaphthalene-8,8'-diol (1) and (2) and its derivatives as new chiral sources. 8,9 The excellent properties of 3 as a proton source for enantioselective protonation of enolates have been explored. 8 These compounds are also expected to function as host molecules that discriminate chirality of organic guest molecules because two hydrogen donors and/or acceptors are located in a highly asymmetric micro environment. In this paper, we report preliminary studies on chiral recognition by 1 and 2 of amino acid derivatives, amino alcohols, and amines caused. 10

¹H-NMR binding studies ¹¹ at 20 °C showed that amines $4 \sim 13$ form diastereomeric complexes with enantiomers 1 (R) and 2 (S) in organic solvents. Table 1 gives the association constants, Ka, and the difference in stability between the diastereomeric complexes, $\Delta\Delta G^{\circ}$. Chiral recognition is most pronounced for the complexes of valine derivative $4 \{\Delta\Delta G^{\circ} = 1.0 \sim 1.2 \text{ kcal/mol} \text{ (enantioselectivity of } \sim 88:12), \text{ entries 1 and 2}.$ The corresponding methyl ester 5 and a free amine 6 showed much reduced enantioselectivity in the binding with 1 and 2 ($\Delta\Delta G^{\circ} \sim 0.3 \text{ kcal/mol}, \text{ entries 3-5}).$ In phenylalanine derivatives 7 ~ 9, chirality was

scarcely recognized by 1 and 2 ($\Delta\Delta G^{\circ} \sim 0.2$ kcal/mol, entries 6-10). ¹² In both amino acid derivatives, zwitterions 4 and 7 caused stronger binding than the corresponding esters 5 and 8, respectively (entries 1 vs 3, 2 vs 4, 6 vs 7). Computer-assisted modeling of the binding motif of 4 and 1 was carried out by a MacroModel/MCMM conformational search¹³ using AMBER*¹⁴ force field. The most stable structure of the complex between 4 and 1 is shown in Figure 1. Characteristic features are: 1) the carboxylate moiety of 4 is hydrogen-bonded with two phenolic OHs,¹⁵ 2) the ammonium moiety of 4 is closely located on the naphthalene ring through ammonium- π interaction, ¹⁶ 3) the *iso*-propyl group of 4 is located closely on the naphthalene ring, which could be ascribed to CH- π interaction¹⁷ or the dispersion force. These calculation results are consistent with the observed ¹H-NMR phenomena in CDCl₃-C₆D₆ (2:1). The protons,

Table 1. Association Constants Ka between Guest Compounds (4~13) and Host Compounds (1 and 2) and the Difference in Binding Energy $\Delta \Delta G^{\circ}$ between Diastereomeric Complexes (T=293K).

entry	guest		Ka (L/mol)		
		solvent	1 (host) 2	∆∆G°a (kcal/mol)
1	4	CDCl ₃	80	11	1.2
2	4	$CDCl_3 - C_6D_6(2:1)$	150	26	1.0
3	5	CDCl ₃	~0	~0	-
4	5	C_6D_6	10	5.6	0.3
5	6	C_6D_6	10	9.8	<0.1
6	7	CDCl ₃	41	37	<0.1
7	8	CDCl ₃	~0	~0	-
8	8	C_6D_6	6.0	5.0	0.1
9	9	CDCl ₃	~0	~0	-
10	9	C_6D_6	15	11	0.2
11	10	CDCl ₃	8.4	11^b	0.2
12	10	C_6D_6	41	34 ^b	0.1
13	11	CDCl ₃	15	15^{c}	<0.1
14	11	C_6D_6	40	35	<0.1
15	12	CDCl ₃	19	19	< 0.1
16	12	C_6D_6	77	78	< 0.1
17	13	CDCl ₃	4.0	2.7^{d}	0.2
18	13	C_6D_6	34	32^d	<0.1

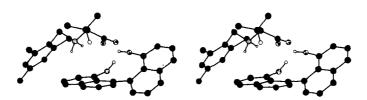
a) Differnce in stability between diatereomeric complexes, guest-1 and guest-2.

$$CO_2R$$
 CO_2Me
 NH
 NH
 NH_2
 N

b) Ka between ent-10 and 1. c) Ka between ent-11 and 1. d) Ka between ent-13 and 1.

-NHC $_{12}$ Ar and -CH(C $_{13}$)2 of 4 (0.014 M), exhibited up-field shifts of 0.41 and 0.27 ppm, respectively, in the presence of 1 (0.037 M). Thus, a three-point binding motif can be assumed; nonetheless, the structure of the host molecule 1 is quite simple. Calculations of the most stable structure 18 of the complex between 4 and 2 indicated a lower stability than that between 4 and 1 by 0.6 kcal/mol, which is roughly consistent with the experimental results shown in Table 1, entries 1 and 2. We should note, however, the above discussions are significant only in the binding of valine derivative 4, since the corresponding phenylalanine derivative 7 showed much reduced enantioselectivity in the binding with 1 and 2. Amino alcohols 10, 11 and amines 12, 13 showed complex formation with 1 and 2 without noticeable chiral recognition (entries $11 \sim 18$). Binding between guests and hosts is stronger in C_6D_6 than in CDCl₃. This implies that the major binding force is hydrogen bonding.

Figure 1



Computer-assisted binding motif (stereoview) between 4 and 1. Hydrogens, except the interacting ones, are omitted for clarity.

We next examined optical resolution through clathrate formation with 2. A mixture of racemic pyrrolidine-2-methanol (14) (110 mg, 1.1 mmol) and 2 (280 mg, 0.89 mmol) in benzene was kept at room temperature to give precipitates, which were then recrystallized twice from benzene to furnish a 1:1 complex of 11 (S, 98% ee) and 2 (153 mg, 36% yield). The absolute configuration and the ee were determined by its transformation into 15.19 The highly diastereoselective clathrate formation does not seem to originate in the diastereoselectivity of the complex formation between 14 and 2 in solution. In both CDCl₃ and C_6D_6 solution, 11 showed complex formation with 1 and 2, but without noticeable chiral recognition (Table 1, entries 13 and 14).

In conclusion, we have shown the potential ability of 1,1'-binaphthalene-8,8'-diol for chiral recognition. Since the simple structure already possesses the functionalities responsible for hydrogen bonding, XH- π , CH- π , and π - π interactions, highly enantioselective recognition would be feasible through further functionalization of the structure.

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