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Structure Study of Host-Guest Molecular Association in Solution and in the Solid State

Fumio Toda,** Koichi Tanaka,* Minoru Ootani,* Atsuhiro Hayashi,* Ikuko Miyahara* and Ken Hirotsu**

Department of Applied Chemistry, Faculty of Engineering, Ehime University, Matsuyama, Ehime 790, Japan
Department of Chemistry, Faculty of Science, Osaka City University, Sumiyoshi-ku, Osaka 558, Japan

A correlation between the mechanism of ¹H NMR shift of a chiral guest by association with an optically active host compound in solution and in the solid state is revealed by an X-ray analysis of the host–guest inclusion crystal.

Optically active hosts, such as 1,6-bis(o-chlorophenyl)-1,6diphenylhexa-2,4-diyne-1,6-diol $1,^{1,2}$ 2,2'-dihydroxy-1,1'binaphthyl $2,^{1,2}$ 4,4',6,6'-tetrachloro-2,2'-bis(hydroxydiphenylmethyl)biphenyl $3,^3$ and tartaric acid derived *trans*bis(hydroxydiphenylmethyl)-2,2-dimethyl-1,3-dioxacyclopentane **4a** and its derivatives **4b** and **4c**^{4,5} can act as NMR chiral shift reagents for determining enantiomeric purity and absolute configuration of various chiral compounds.

For example, the methine proton signal of racemic-1phenylethylamine **5a**, which appears as a quartet centred at δ 4.25 in the ¹H NMR spectrum[†] is split into two quartets centred at δ 3.63 and 3.60 in the presence of 2 equiv. of **4a**. The two signals centred at δ 3.60 and 3.63 were assigned to the methine proton of (S)-(-)-**5b** and the (R)-(+)-enantiomer **5c**, respectively, by measurement of the ¹H NMR spectra of the optically pure **5b** and **5c** in the presence of **4a**.

The above data can be interpreted by assuming the following molecular association of 4a and 5 in CDCl₃. In a molecular complex of 4a and 5b produced by their association through hydrogen-bond formation between the hydroxy group of 4a and the amine nitrogen atom of 5b, the methine proton of 5b is located in the position that is shielded by a phenyl ring of 4a and 5c, however, the methine proton of 5c is

located in the position that is deshielded and is shifted up-field. In order to confirm the assumption, host-guest inclusion crystal of **4a** and **5b** was prepared and its X-ray structure was analysed.

When a solution of 4a⁶ (5 g, 10.7 mmol) and 5a (1.3 g, 10.7 mmol) in toluene-hexane (1:2, 30 ml) was kept at room temp. for 12 h, a 2:1 inclusion complex of 4a and 5b of 91% e.e. (enantiomeric excess) was obtained as colourless prisms (3.5 g, 62% yield). Recrystallisation of the inclusion crystal from toluene gave the complex 6 of pure enantiomer (3 g, 52% yield, m.p. is not clear), which upon heating *in vacuo* gave 5b of 100% e.e. by distillation {0.32 g, 49% yield, $[\alpha]_D^{25}$ -22.5 (*c* 0.18, MeOH)}. The efficient optical resolution of the guest compound by inclusion complexation with optically active host compound has thus been established.⁷

The X-ray crystal structure of 6^{\dagger} shows that two host molecules are related by an approximate twofold axis passing through the midpoints between O(3) and O(3') and O(4) and O(4'). All the hydrophilic groups are concentrated in the interior of the complex to make a cyclic hydrogen-bond system [...O(4)-H...N(1)-H...O(4')-H...O(3')-H...O(3)-H...] with one of the two N-H groups of **5b** free from a hydrogen bond, while the exterior of the complex is hydrophobic and no inter-complex hydrogen bond is formed. The host-guest short contacts are mainly found between the phenyl group [C(26), C(27),...C(31)] on C(7) of **4a** on the left in Fig. 1 and the -CH(NH₂)Me part of **5b**, with the methine group of **5b** shielded by the phenyl group.

 $^{^{\}dagger}$ All ^{1}H NMR spectra were measured in CDCl3 at room temp. by using a 60 MHz JEOL PMX-60SI spectrometer.

Similarly, 4b also acts as a chiral shift reagent. For example, the methyl ¹H signal of racemic 2-methylpiperidine 7a, which appears as a doublet centred at δ 1.09, is split into two doublets centred at δ 0.82 and 0.87 in the presence of an equimolar amount of 4b. The two signals centred at δ 0.82 and 0.87 were assigned to the methyl proton of (R)-(-)-7b and (S)-(+)-enantiomer 7c respectively, by measurement of the ¹H NMR spectra of the optically pure 7b and 7c in the



C(20)

C(21

O(4)

N(1)

C(11) C(12)

2(10

C(15)

C(34

C(13)

C(35)

C(38)

C(14)

_C(36)

presence of 4b. The splitting of the methyl proton signal of 7a in the presence of 4b can also be interpreted as a host-guest molecular association in CDCl₃. In a molecular complex of 4b

C(18)

C(19)



Fig. 1 ORTEP drawing of host-guest (2:1) complex of 4a and 5b. Hydrogen bonds are shown by dotted lines. Hydrogen bond distances (Å): O(4)-N(1) 2.730(7); N(1)-O(4') 2.955(7); O(4')-O(3') 2.652(5); O(3')-O(3) 2.860(5); O(3)-O(4) 2.677(4).

and 7b, the methyl proton of 7b is located in the position shielded by a phenyl ring of 4b and is shifted downfield. In a molecular complex of 4b and 7c, however, the methyl proton of 7c is located in the position deshielded and is shifted up-field. In order to confirm the assumption, an inclusion crystal of 4b and 7c was prepared and its structure was studied by X-ray analysis.

When a solution of **4b** (5 g, 10.2 mmol) and **7a** (2 g, 20.2 mmol) in toluene-hexane (1:1, 20 ml) was kept at room temp. for 3 h, a 1:1 inclusion complex of **4b** and **7c** of 42% e.e. was formed as colourless needles (5.4 g, 90% yield). Three recrystallisations of the crude complex from toluene gave the complex **8** of pure enantiomer (1.6 g, 30% yield, m.p. is not clear), which upon heating *in vacuo* gave **7c** of 100% e.e {0.26 g, 26% yield, $[\alpha]_D^{25} + 4.5$ (c 0.85, MeOH)}.

The X-ray crystal structure analysis of $8\ddagger$ showed that the hydrophilic groups of 4b and 7c are suited to make a hydrogenbond network [O(3)-H···O(4)-H···N(1)]. Similarly to the complex of 4a and 5b, 6, the exterior of the complex of 4b and 7c, 8 is mostly hydrophobic with no inter-complex hydrogen bond (NH of 7c does not take part in the hydrogen bond). The phenyl ring [C(28), C(29), ···C(33)] on C(9) of 4b interacts

Crystal data for 8: $C_{33}H_{32}O_4C_6H_{13}N M = 591.76$, orthorhombic, space group $P2_12_12_1$, a = 22.796(4), b = 15.436(2), c = 9.298(3) Å, U = 3272(1) Å³, Z = 4, $D_c = 1.20$ g cm⁻³, $\mu = 0.71$ cm⁻¹. 3261 Reflections with $2\theta < 50.0^{\circ}$ were recorded on a Rigaku AFC-6A four-circle diffractometer using graphite-monochromated Mo-K α radiation. Of these, 2404 with $F > 3\sigma(F)$ were judged as observed. The structure was solved using SHELX-86.9 Full-matrix least-squares refinement with anisotropic temperature to R = 0.045 and $R_w =$ $0.074.^{10}$ For both structures, atomic coordinates, bond distances and angles, and thermal parameters have been deposited at the Cambridge Crystallographic Data Centre. See Notice to Authors, Issue No. 1. mainly with N(1)-C(38) moiety of 7c and no short contact is found between the methyl group of 7c and the phenyl rings of 4b.

The interpretation of the ¹H NMR spectra by assuming the molecular association described above is supported by the X-ray structures of these two complexes, 6 and 8. Interestingly, the relative orientation of 4a on the left and 5b in Fig. 1 resembles that of 4b and 7c in Fig. 2. Moreover, the same type of host-guest arrangements have been found in the 1:1 inclusion complex crystal of 4a with (R)-(-)-6-methyl-[4.4.0]dec-1-en-3-one and with (R)-(-)-methylbicyclo-[4.4.0]dec-1-ene-3,7-dione.⁸ Thus, these results may suggest that the host-guest association of 4b with 7c and 4a (on the left in Fig. 2) with 5b may reflect approximately host-guest interactions in CDCl₃. An important feature of the host-guest association in this orientation is that the complex is stabilised by a hydrogen-bond formation of a hydroxy group of a host molecules as a donnor with N atom of a guest molecule as an acceptor, and the van der Waals interactions between the phenyl rings of a host molecule and a guest molecule.

We thank the Ministry of Education, Science and Culture, Japan, for a Grant-in-Aid for Cooperative Research (A), No. 03303002 and for Scientific Research (B), No. 04453102.

Received, 14th May 1993; Com. 3/02734A

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[‡] Crystal data for 6: 2(C₃₁H₃₀O₄) C₈H₁₁N M = 1054.28, orthorhombic, space group $P_{21}2_{12}1$, a = 20.308(6), b = 17.173(2), c = 17.042(2) Å, U = 5943(2) Å³, Z = 4, $D_c = 1.17$ g cm⁻³, $\mu = 0.71$ cm⁻¹, 5780 Reflections with $2\theta < 50.0^{\circ}$ were recorded on a Rigaku AFC-6A four-circle diffractometer using graphite monochromated Mo-Ka radiation. Of these, 3566 with $F > 3\sigma(F)$ were judged as observed. The structure was solved using SHELX-86.⁹ The hydrogen atoms of the methyl group on C(32) could not be located because of large thermal motion. Full-matrix least-squares refinement with anisotropic temperature factors for non-hydrogen atoms and isotropic hydrogens converged to R = 0.057 and $R_w = 0.084.^{10}$