

X-Ray Structural Study of a 1:2:1 Inclusion Complex of Non-centrosymmetric
1,6,9,14-Tetrahydroxy-1,6,9,14-tetra-*t*-butylcyclohexadeca-
2,4,7,10,12,15-hexayne, Ethanol, and Water

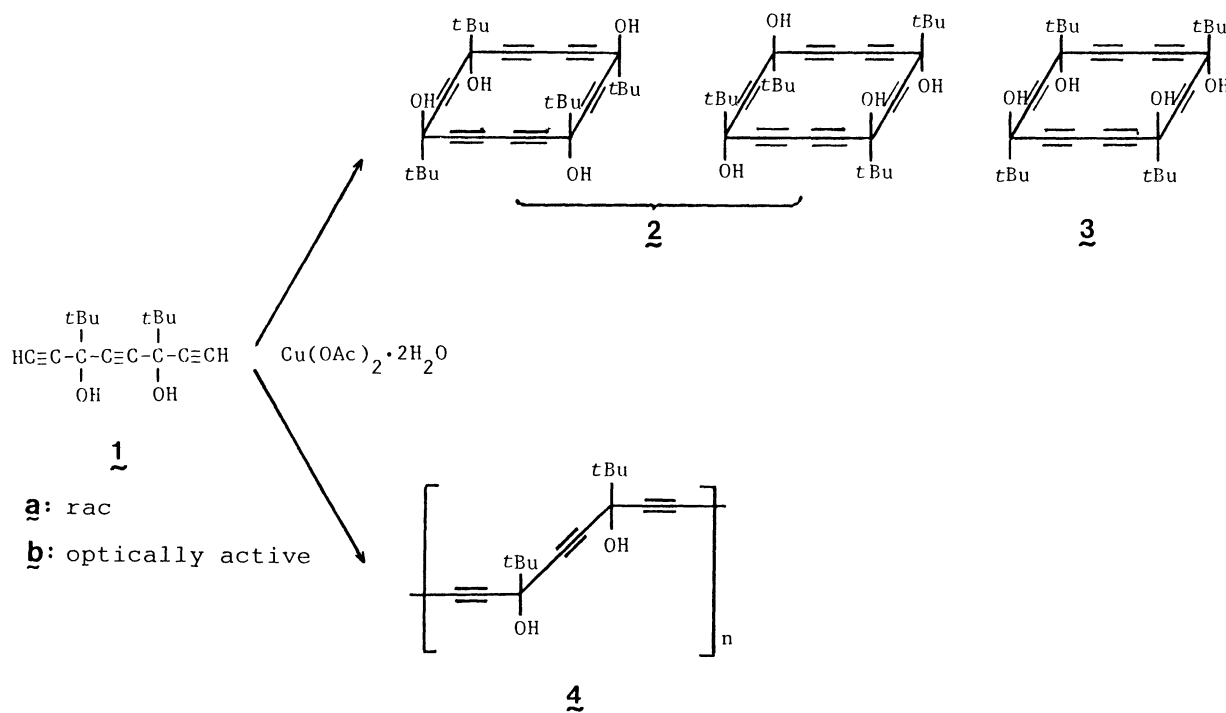
Fumio TODA,^{*} Kōji OKADA, and Thomas C.W. MAK^{*†}

Department of Industrial Chemistry, Faculty of Engineering,
Ehime University, Matsuyama 790

[†]Department of Chemistry, The Chinese University of Hong Kong,
Shatin, New Territories, Hong Kong

The structure of the title cyclic hexayne has been elucidated by X-ray analysis of its 1:2:1 inclusion complex with EtOH and H₂O. The important role of H₂O in consolidating the disordered, hydrogen-bonded host lattice is also clarified.

Previously we reported that oxidative coupling of racemic (1a) and optically active (1b) 3,6-di-*t*-butylocta-1,4,7-triyn-3,6-diol gives cyclic dimeric (2 or 3) and optically-active polymeric (4) products, respectively.¹⁾ However, the cyclic dimer was not firmly established as either the racemic (2) or the *meso* (3) compound.



Recrystallization of (2) (white powder, mp 300 °C)¹⁾ from absolute ethanol did not give crystals of good quality. However, use of aqueous ethanol as a solvent yielded colorless prisms (mp not sharp) of the title complex (5), whose stoichiometry and structure were then elucidated by X-ray crystallography. The present study has shown the cyclic hexayne to be the non-centrosymmetric isomer (2) and (5) to be an inclusion compound.

A selected crystal of 5 with dimensions of 0.40x0.40x0.40 mm³ was sealed inside a 0.5 mm Lindemann glass capillary, and intensity data were collected on a Nicolet R3m/V diffractometer at 22 °C using Mo-K α radiation (λ = 0.71073 Å) according to established procedures.²⁾

Crystal data: 5, C₃₂H₄₀O₄·2C₂H₅OH·H₂O, FW = 598.81, tetragonal, space group P4₁n2 (No. 118), a = 10.464(3), c = 17.153(6) Å, V = 1878(1) Å³, Z = 2, $F(000)$ = 652, D_m (floatation in KI/H₂O) = 1.067, D_c = 1.059 g cm⁻³, μ = 0.41 cm⁻¹, $2\theta_{max}$ = 50°, 1681 unique reflections. The structure was solved by direct phase determination based on random starts and negative quartets.^{3,4)} The cyclic hexayne molecule (Fig. 1) is located at Wyckoff position 2(c) of site symmetry 222, and the water molecule O2 occupies position 2(a) of site symmetry $\bar{4}$. Half site-occupancy and disorder of the ethanol molecule, which occupies a general position, became evident in the course of refinement. In the model adopted, the ethanol molecule is represented by atoms O3[$\frac{1}{2}$], C9[$\frac{1}{2}$], C10[$\frac{1}{2}$] and C11[$\frac{1}{2}$] with site occupancy factors enclosed in square brackets. Atoms C9, C10, and C11 were varied isotropically, and all other non-hydrogen atoms anisotropically. The methyl H atoms in the cyclic hexayne were generated geometrically (C-H fixed at 0.96 Å), and the hydroxyl H(O1) atom was located from a difference map. These H atoms were included in structure-factor calculations in least-squares cycles with assigned isotropic temperature factors. Refinement of 1403 observed data [$|F_o| > 3\sigma(|F_o|)$] on 114 parameters using the weighting scheme $w = [\sigma^2(|F_o|) + 0.0025|F_o|^2]^{-1}$ converged to R_F = 0.089. All computations were performed on a DEC MicroVAX-II computer with the SHELXTL PLUS package.³⁻⁶⁾

The measured dimensions of the cyclic dimer of D₂ molecular symmetry is given in the legend of Fig. 1. The C₂ axis passing through the mid-points of the C4-C4^b and C4^a-C4^c bonds runs parallel to the crystallographic c axis. Another C₂ axis passes through the centers of the C2-C2^a and C2^b-C2^c bonds, whereas the third is normal to the mean plane of the molecule. The central sixteen-membered ring is distinctly non-planar, and its edges are distorted from linearity in a barrel-like manner, as shown by the values of the relevant bond angles and torsion angles in Fig. 1. In accordance with expectation, the unconjugated C \equiv C bond is found to be significantly shorter than the conjugated one, and the *t*-butyl group is in a staggered arrangement with respect to the ring atoms.

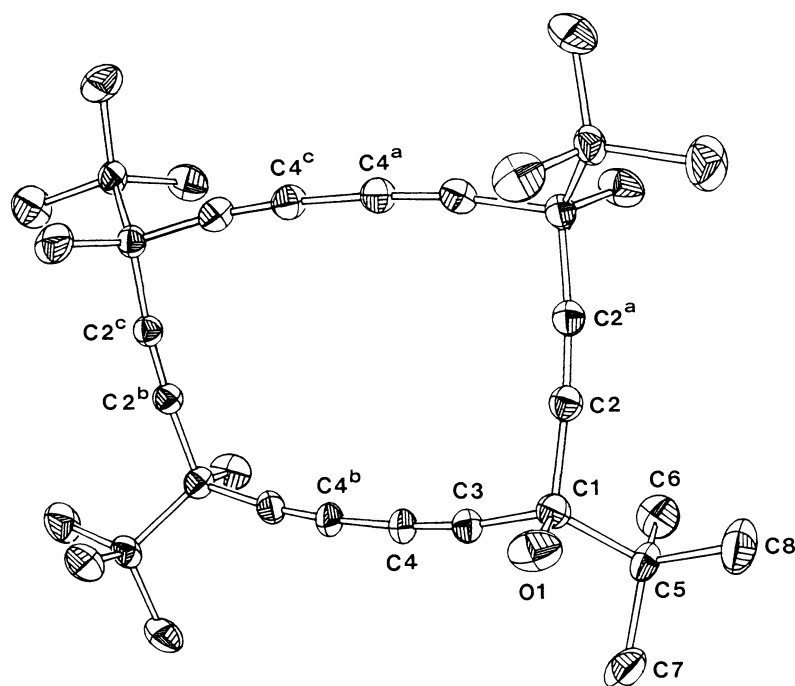


Fig. 1. The structure of the cyclic hexayne molecule with atom labelling. Thermal ellipsoids are drawn at the 30% probability level. Selected molecular dimensions (bond lengths in Å, angles in degrees, estimated standard deviations in parentheses): C1-C2 1.472(6), C2-C2^a 1.168(6), C1-C3 1.494(6), C3-C4 1.184(6), C4-C4^b 1.381(6), C1-C5 1.571(6), C1-O1 1.429(5); C1-C2-C2^a 173.8(3), C1-C3-C4 172.3(3), C3-C4-C4^b 175.3(3), C2-C1-C3 105.9(3), C2-C1-C5 111.6(3), C3-C1-C5 111.1(3), C2-C1-O1 108.6(4), C3-C1-O1 109.8(3), C5-C1-O1 109.7(5); torsion angles C1-C2-C2^a-C1^a 21.1(4), C1-C3-C4-C4^b -24.3(4), C3-C4-C4^b-C3^b 20.8(4), O1-C1-C5-C6 -178.0(5). Symmetry code: ^a $\frac{1}{2}-y, \frac{1}{2}-x, \frac{1}{2}-z$; ^b $-\underline{x}, 1-\underline{y}, \underline{z}$; ^c $-\frac{1}{2}+\underline{y}, \frac{1}{2}+\underline{x}, \frac{1}{2}-\underline{z}$.

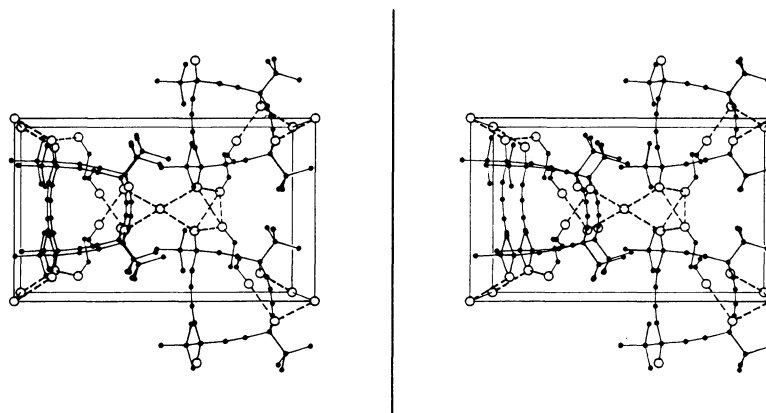


Fig. 2. Stereodrawing of the crystal structure of **5**. The origin of the unit cell lies at the upper left corner, with a pointing toward the reader, b vertically downward, and c from left to right. Broken lines represent O...O hydrogen bonds. Note that the ethanol molecule is two-fold disordered, and one of the possible methyl carbon positions (atom C11) has been omitted for clarity.

A stereo view of the crystal packing is illustrated in Fig. 2. The water molecule forms hydrogen bonds [$O1...O2 = 2.806(5) \text{ \AA}$] with the hydroxyl groups of four neighboring hexayne molecules. The ethanol molecule is disordered over two sites, and its methyl group furthermore exhibits two-fold positional disorder. The O atoms of the hexayne and disordered ethanol molecules are also linked by hydrogen bonds [$O1...O3^a = 2.662(8)$, $O1...O3(\frac{1}{2}+\underline{y}, \frac{1}{2}+\underline{x}, \frac{1}{2}-\underline{z}) = 2.843(8) \text{ \AA}$]. Compound 5 can thus be considered as an inclusion complex comprising a host lattice built of hexayne and water molecules which accommodates the ethanol guest species. It is conceivable that two 'half' ethanol molecules related by a crystallographic C_2 axis may be replaced by another guest species of similar bulk and hydrogen-bonding capability, such as 1,2-ethanediol, to form an isomorphous inclusion complex. When the guest molecule fails to satisfy these requirements, the disordered hydrogen-bond system of the present host lattice will most likely be disrupted, and the complex is expected to crystallize in a space group of lower symmetry.

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