1,6,7,10-Tetramethylfluoranthene: Synthesis and Structure of a Twisted Polynuclear Aromatic Hydrocarbon.

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Abstract: The structure and synthesis of the overcrowded polynuclear aromatic 1,6,7,10tetramethylfluoranthene, 1, is presented. The structures of 1 and 11 have been determined by X-ray diffraction to be twisted into conformations of approximately C_2 symmetry due to the buttressing of the flanking methyl groups. Semi-empirical calculations (AM1) on 1 find two minima, twist and fold; the twist is of lower energy. The energy for enantiomerization of 1 is set at below 7.0 kcal/mol by variable temperature NMR. Computations predict a likely path to account for the ribbon twist.

Steric crowding in aromatic structures typically results in twisting² or folding³ out-of-plane distortion of the molecule.⁴ Synthesis and elucidation of the stereochemistry of such systems occupies an important position in the chemistry of polynuclear aromatic hydrocarbons.^{5,6} Stereochemical analysis predicts 1,6,7,10-tetramethylfluoranthene (1) to be a helically-twisted, strained, aromatic molecule.⁷ This report presents the synthesis and structural characterization of 1 and related compounds.⁸

Modification of an earlier scheme allowed us to prepare 2,7-dimethylacenaphthenequinone⁹ (4) from 2,7-dimethylnaphthalene¹⁰ (3) in 6 steps (Scheme 1). Condensation of 4 with 3-pentanone and base formed the carbinol 5. Reaction of 5 with a variety of acetylene equivalents resulted in fluoranthenes 1, 6, and 7. The diester 6 was further derivatized by reduction with lithium aluminum hydride selectively to the diol 9 or exhaustively to 1,6,7,8,9,10-hexamethylfluoranthene (8). Treatment of 1 with NBS in carbon tetrachloride yielded 11.

The X-ray structures of 1, 6, and 11 show a helical twist in the aromatic ribbon.^{11,12} The twist is nearly continuous throughout the molecule. The dihedral angle between the best plane through the benzene ring and the best plane through the naphthalene ring is 16 and 18°, respectively, and the dihedral angle between the top bond of the benzene and the vector across the free peri positions in the naphthalene is 19 and 23°, respectively. The proximal $C(sp^3)$ to $C(sp^3)$ distance in 1 is 3.09 Å and in 11 is 3.25 Å; both fall within the sum of van der Waals (vdW) distance of 4.0 Å. The bromine to bromine distance in 11 is 3.9 Å, close to the sum of vdW radii (Figure 1).

The dynamic stereochemistry of this system in solution was probed through the diacetate 10. The methylene protons served as a probe of the chirotopicity of the methylene group as a whole. They give rise to an AA' pattern in the fluxional molecule (the achirotopic state), and an AB pattern in the static molecule (the chirotopic state). At -100 °C in deuterofreon¹³ the methylene protons split out. A dynamic process of ca. 7.0 ± 0.5 kcal/ mol was found.¹⁴ On the basis of our previous work on hexasubstituted benzenes,¹⁵ we assigned this



Scheme 1. a) Paraformaldehyde, CH₃CO₂H/HCl, 50 °C, 24 h, 62%; b) KCN, acetone/H₂O, reflux, 12 h, 99%; c) H₂SO₄, CH₃CO₂H, H₂O, reflux, 6 h, 90%; d) SOCl₂, CHCl₃, reflux, 1hr; e) AlCl₃, nitrobenzene, rt, 48 h, 82% (two steps); f) SeO₂, dioxane/H₂O, 60 °C, 48 h; g) 3-pentanone, MeOH, KOH, rt, 30 min; h) Norbornadiene, acetic anhydride, 125 °C, 30 h; i) Maleic anhydride, acetic anhydride; (j) Dimethyl acetylenedicarboxylate, acetic anhydride. k) LiAlH₄, THF. l) Acetyl chloride. m) NBS (5 eq.), CCl₄, reflux, 48h.

to the rotation about the C(aromatic)-C(acetoxymethyl) bond and not to the twist interconversion. Nonetheless, this sets an upper limit for the barrier for reversal of the ribbon twist.



Figure 1. X-ray structures of 1 (right) and 11 (left) showing the twisted (15-20°) aromatic ribbon and juxtaposed methyl groups (3.1-3.3 Å). Picture from MacMoMo, M. Dobler (ETH-Zürich).

Semi-empirical quantum calculations using the AM1 Hamiltonian¹⁶ predict two low energy conformations for 1, a twist and a fold, with the twist lying ca. 1.2 kcal/mol (2.7 kcal/mol: PCMODEL) below the fold. The geometry of the twist compares favorably with the X-ray data: the twist angle is ca. 15° (1, 16°; 11, 18°: X-ray), and the methyl-to-methyl distance is 3.05 Å (1, 3.09 Å; 11, 3.25 Å: X-ray). The flat conformation lies ca. 15 kcal/mol higher in energy than the twist. From our dynamic NMR work and calculations on similar systems, we can exclude the flat structure as a point on the path to ribbon reversal. The most likely path is twist-fold-twist by way of a transition structure in which only one set of proximal methyl groups pass one another.

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