Supplementary data for this paper are available from the IUCr electronic archives (Reference: DA1077). Services for accessing these data are described at the back of the journal.

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Acta Cryst. (1999). C55, 2090-2091

2-tert-Butyl-9,10-dibromoanthracene

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(Received 9 June 1999; accepted 17 September 1999)

Abstract

The title compound, $C_{18}H_{16}Br_2$, crystallized on a crystallographic mirror plane. There was disorder of the 'Bu group with a major orientation at 59%. Interplanar spacings are 3.509 (4) Å.

Comment

The title compound, (I), was structurally characterized in order to measure interplanar distances for studies aimed at assessing the forces of attraction in this molecule, and to make comparisons with other molecules such as anthracene (Lehmann & Pawley, 1972) and 9,10dibromoanthracene (Kitaigorodski, 1946; Trotter, 1958, 1986; Bulgarovskaya et al., 1987) of a similar nature. We find that the interplanar spacing in (I) of 3.509(4) Å is similar to that in 9,10-dibromoanthracene of 3.52(1) Å (Trotter, 1986). It is also interesting to note that the packing arrangement in 9,10-dibromoanthracene, where the molecules are stacked on top of each other, is very different from that observed in (I). Clearly, the 'Bu group is responsible for these differences. Additionally, the closest $Br \cdots Br$ intermolecular distance of 4.004 (2) Å in (I) is smaller than that of 4.068 (1) Å in 9,10-dibromoanthracene (Trotter, 1968). It is likely that the skewed arrangement of the rings in (I) which reduces Br-Br repulsions, also results in a decrease in the interplanar spacings in (I).



Molecules of (I) in the crystal are arranged in a herring-bone manner parallel to the *bc* plane and perpendicular to the *a* axis. The angle for the intramolecular $Br \cdots Br$ vectors situated between adjacent coplanar rows is 90.9 (1)°. Adjacent pairs of molecules are skewed, with 'Bu groups on opposite sides and displaced by one ring so that only the two fused rings furthest from the 'Bu group are aligned on top of each other. The bond distance and angles within the anthracene rings in (I) and 9,10-dibromoanthracene are similar.



Fig. 1. View of the major contributor to (I) (50% probability displacement ellipsoids). H atoms, represented by circles of arbitrary radii, are not labeled.

Experimental

Crystals of (I) prepared as detailed previously (Lee & Mendenhall, 1988) were obtained by slow crystallization from an ethanol solution.

Crystal data

$C_{18}H_{16}Br_2$	Mo $K\alpha$ radiation
$M_r = 392.14$	$\lambda = 0.71069 \text{ Å}$
Orthorhombic	Cell parameters from 25
Pnma	reflections
a = 23.433(5)Å	$\theta = 10.4 - 15.0^{\circ}$
b = 7.018 (4) Å	$\mu = 5.15 \text{ mm}^{-1}$
c = 9.554 (4) Å	T = 293 (2) K
$V = 1571.2 (12) \text{ Å}^3$	Prism
Z = 4	0.22 \times 0.20 \times 0.08 mm
$D_x = 1.658 \text{ Mg m}^{-3}$	Yellow
D_m not measured	

Data collection 1183 reflections with Enraf-Nonius CAD-4 diffractometer $I > 2\sigma(I)$ $\theta_{\rm max} = 27.47^{\circ}$ Non-profiled ω scans $h=0\to 30$ Absorption correction: $k = 0 \rightarrow 9$ ψ scan (North *et al.*, $l=0 \rightarrow 12$ 1968) 3 standard reflections $T_{\rm min} = 0.304, T_{\rm max} = 0.674$ 1930 measured reflections frequency: 60 min 1930 independent reflections intensity decay: 2%

Refinement

Refinement on F^2	$(\Delta/\sigma)_{\rm max} = 0.002$
R(F) = 0.040	$\Delta \rho_{\rm max} = 0.73 \ {\rm e} \ {\rm \AA}^{-3}$
$wR(F^2) = 0.112$	$\Delta \rho_{\rm min} = -0.44 \ {\rm e} \ {\rm \AA}^{-3}$
S = 1.028	Extinction correction:
1930 reflections	SHELXL97 (Sheldrick,
135 parameters	1997)
H atoms constrained	Extinction coefficient:
$w = 1/[\sigma^2(F_o^2) + (0.0642P)^2]$	0.0006 (4)
+ 0.26P]	Scattering factors from
where $P = [Max(F_o^2, 0)]$	International Tables for
$+ 2F_c^2$]/3	Crystallography (Vol. C)

Table 1. Selected geometric parameters (Å, °)

Br1—C9	1.905 (5)	C8aC9	1.412 (7)
Br2—C10	1.903 (5)	C9C9a	1.401 (7)
C9a—C9—Br1 C8a—C9—Br1	1.392 (7) 118.2 (4) 119.0 (4)	C4a—C10—Br2 C10a—C10—Br2	118.6 (4) 118.8 (4)

A difference Fourier map clearly indicated two regions of electron density suggestive of disorder for with 'Bu group. C atoms were inserted and the sum of occupancies of the relevant C atoms were constrained to unity and then refinement continued to convergence both on positions and occupancies.

Data collection: CAD-4 EXPRESS (Enraf-Nonius, 1994). Cell refinement: CAD-4 EXPRESS. Data reduction: XCAD4 (Harms & Wocadlo, 1995). Program(s) used to solve structure: SIR97 (Altomare et al., 1999). Program(s) used to refine structure: SHELXL97 (Sheldrick, 1997). Molecular graphics: ORTEP-3 for Windows (Farrugia, 1997). Software used to prepare material for publication: WinGX (Farrugia, 1998).

RLL thanks Michigan Technological University for support and reviewers for helpful comments.

Supplementary data for this paper are available from the IUCr electronic archives (Reference: FR1224). Services for accessing these data are described at the back of the journal.

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The *tert*-butylammonium perchlorate complex of 2,3-naphtho-18-crown-6 at 115 K

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(Received 29 June 1999; accepted 10 August 1999)

Abstract

In the title complex, *tert*-butylammonium perchlorate– 2,5,8,11,14,17-hexaoxatricyclo[16.8.0.0^{20,25}]hexacosa-1(26),18,20(25),21,23-pentaene–ethyl acetate–dichloromethane (4/4/1/1), C₄H₁₂N⁺·C₂₀H₂₆O₆·ClO₄⁻·0.25C₄-H₈O₂·0.25CH₂Cl₂, the *tert*-butylammonium cation binds to the macrocyclic host (Chemical Abstracts name: 2,3,5,6,8,9,11,12,14,15-decahydro-1,4,7,10,13,16-hexaoxanaphtho[2,3-*b*]cyclooctadecin) in the expected tripod arrangement, while the perchlorate anion links naphthyl groups in the crystal through C—H···O—Cl—O···H— C interactions. Thermal motion analysis indicates that the *tert*-butylammonium group and the perchlorate anion each librate with respect to the host, with amplitudes of 6.2 (4) and 11.4 (2)°, respectively.

Comment

The host in the title compound, (I), has been used as a standard in solution studies of structures of hosts and their complexing affinities; see, for example, Kyba *et al.* (1977) and Helgeson *et al.* (1979). The *tert*-butyl-

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