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Naotake Nakamura* and Akira Ohishi

Department of Applied Chemistry, College of Science and Engineering, Ritsumeikan University, 1-1-1 Nojihigashi, Kusatsu, Shiga 525-8577, Japan

Correspondence e-mail: nakamura@se.ritsumei.ac.jp

Key indicators

Single-crystal X-ray study T = 296 KMean σ (C–C) = 0.007 Å R factor = 0.056 wR factor = 0.108 Data-to-parameter ratio = 16.2

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

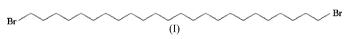
1,24-Dibromotetracosane

In the title compound, $C_{24}H_{48}Br_2$, the molecule is centrosymmetric and the molecular skeleton including both terminal Br atoms has an all-trans conformation. The molecules form layers with a thickness of c/2. These features are similar to the smectic C_A phase of liquid crystals.

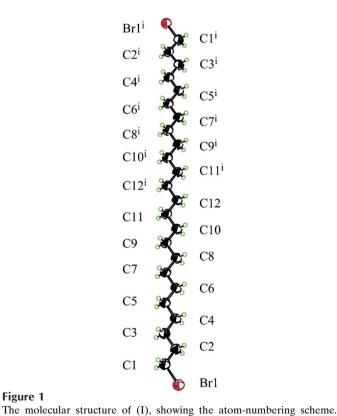
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Comment

Normal long-chain compounds can be regarded as having a rod-like shape, and the molecules in the crystalline state form a layered structure similar to those of the smectic liquid crystalline phase. Moreover, some of these long-chain compounds exhibit a high-temperature rotator phase just below their melting points, in which molecules have motional freedom to some degree, comparable with that in liquid crystals. Thus, these long-chain compounds have been studied as model compounds for smectic liquid crystals.



Many researchers have analyzed the crystal structures of many different kinds of normal long-chain aliphatic

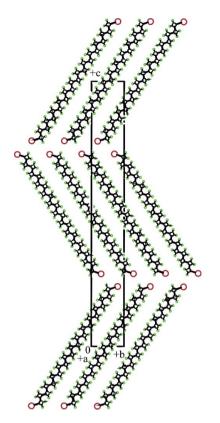


Displacement ellipsoids are drawn at the 50% probability level

Figure 1

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[symmetry code: (i) -x, 1 - y, 2 - z].





compounds. Recently, we have systematically analysed the crystal structures of the alkane- α , ω -diols containing 10–19 and 21–23 C atoms (Nakamura *et al.*, 2001*a*; Uno *et al.*, 2002) and the alkane- α , ω -dithiols containing 12, 19–21 and 23 C atoms (Nakamura *et al.*, 2001*b*,*c*, 2004, 2005*a*,*b*). In the present paper, we report the results of the crystal structure analysis of the title compound, (I), in order to elucidate the effect of the terminal groups in normal long-chain compounds in constructing a layer structure.

The molecular structure of (I) is shown in Fig. 1. The molecule is centrosymmetric and all torsion angles are close to $\pm 180^{\circ}$, that is, the molecular structure including both terminal Br atoms has an all-trans conformation. The layers are arranged in a zigzag manner between the neighboring layers, forming a herringbone motif similar to the smectic C_A phase of liquid crystals, as shown in Fig. 2. The layers are stacked closely in such a way that the α -CH₂ groups are allowed to fit into the grooves formed by Br atoms, with the nearest contact $Br1 \cdot \cdot \cdot Br1^{i} = 3.754$ (2) Å [symmetry code: (i) $4 - x, y - \frac{1}{2}, \frac{3}{2} - \frac{1}{2}$ z], agreeing closely with the van der Waals contact of 3.75 Å (Rowland & Taylor, 1996). Such a close packing is observed in the even-numbered alkane- α, ω -diols containing 4–19 and 21– 23 C atoms (Thalladi *et al.*, 2000) and the α,ω -dichloroalkanes containing 16, 20 and 26 C atoms (Takamizawa et al., 1992; Nakamura & Shimizu, 2004). Fig. 3 shows the projection of the crystal structure of (I) along the b axis. The molecules form layers with a thickness of c/2. In the layer, the long axes of the molecules are inclined to the *ab* plane. This layer structure is

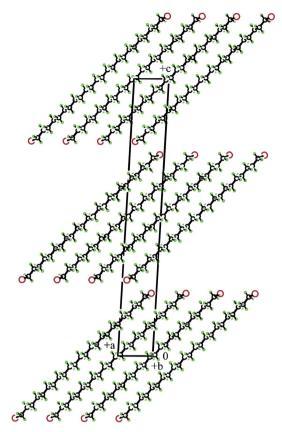


Figure 3 The projection of the crystal structure of (I) along the *b* axis.

similar to that of the triclinic structure of the even-numbered *n*-alkanes containing 6–22C atoms (Nyburg & Gerson, 1992). It is considered that the arrangement in the layer is influenced by the steric and electrostatic repulsion of Br atoms at both ends.

The main features of the structure of (I) are similar to those of the homologous series with an even number of C atoms, *viz*. 1,12-dibromododecane (Kulpe *et al.*, 1981), 1,14-dibromotetradecane (Uno & Nakamura, 2003), 1,16-dibromohexadecane (Kobayashi *et al.*, 1995), 1,18-dibromooctadecane (Nakamura *et al.*, 1993), 1,20-dibromoicosane (Nakamura & Ohishi, 2004) and 1,26-dibromohexacosane (Takamizawa *et al.*, 1992).

Experimental

Thin plate-like crystals of (I) were grown by slow evaporation of a solution in a mixture of ethyl acetate and ethanol (2:1). The well developed face of the crystal is (001).

Crystal	data
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$C_{24}H_{48}Br_2$	$D_x = 1.276 \text{ Mg m}^{-3}$
$M_r = 496.45$	Cu $K\alpha$ radiation
Monoclinic, $P2_1/c$	Cell parameters from 24
a = 5.482 (3) Å	reflections
b = 5.381 (2) Å	$\theta = 9.2 - 17.1^{\circ}$
c = 43.859 (2) Å	$\mu = 3.99 \text{ mm}^{-1}$
$\beta = 93.07 \ (2)^{\circ}$	T = 296 (1) K
$V = 1291.9 (9) \text{ Å}^3$	Plate, colourless
Z = 2	$0.51 \times 0.41 \times 0.01 \text{ mm}$

Data collection

 $R_{\rm int} = 0.040$ Rigaku AFC-5R diffractometer $\theta_{\rm max} = 70.8^{\circ}$ ω scans Absorption correction: numerical $h = -1 \rightarrow 6$ (ABSCOR; Higashi, 1999) $k = -1 \rightarrow 6$ $T_{\min} = 0.175, \ \tilde{T}_{\max} = 0.961$ $l = -53 \rightarrow 53$ 3 standard reflections 4250 measured reflections 2374 independent reflections every 150 reflections 1525 reflections with $F^2 > 2\sigma(F^2)$ intensity decay: 14.0%

Refinement

 Refinement on F^2 $w = 1/[0.0001F_o^2 + 8.45\sigma^2(F_o) + 0.525]/(4F_o^2)$
 $R[F^2 > 2\sigma(F^2)] = 0.056$ $0.525]/(4F_o^2)$
 $wR(F^2) = 0.108$ $(\Delta/\sigma)_{max} < 0.001$

 S = 1.00 $\Delta\rho_{max} = 1.07$ e Å⁻³

 2374 reflections
 $\Delta\rho_{min} = -1.26$ e Å⁻³

 143 parameters
 Extinction correction: Larson

 All H-atom parameters refined
 (1970)

Table 1Selected geometric parameters (Å, °).

Br1-C1	1.926 (5)	C7-C8	1.520 (6)
C1-C2	1.493 (6)	C8-C9	1.518 (6)
C2-C3	1.506 (7)	C9-C10	1.521 (6)
C3-C4	1.520 (6)	C10-C11	1.509 (6)
C4-C5	1.514 (6)	C11-C12	1.526 (6)
C5-C6	1.526 (6)	C12-C12 ⁱ	1.506 (8)
C6-C7	1.516 (6)		
Br1-C1-C2-C3	179.3 (3)	C6-C7-C8-C9	-179.5 (4)
C1-C2-C3-C4	-179.9(5)	C7-C8-C9-C10	180.0 (4)
C2-C3-C4-C5	-179.2(4)	C8-C9-C10-C11	-179.8(4)
C3-C4-C5-C6	180.0 (4)	C9-C10-C11-C12	-179.8(4)
C4-C5-C6-C7	-179.2(4)	C10-C11-C12-C12i	-179.7 (4)
C5-C6-C7-C8	-179.3 (4)	$C11 - C12 - C12^i - C11^i$	180.0 (4)

Symmetry code: (i) -x, -y + 1, -z + 2.

H atoms were positioned geometrically and treated as riding, with C-H distances of 0.95 Å and $U_{\rm iso}(\rm H) = 1.2 U_{eq}(\rm parent)$. The intensity decay was somewhat high (14.0%) because a longer time was needed to collect the reflection data.

Data collection: *MSC/AFC Diffractometer Control Software* (Molecular Structure Corporation, 1992); cell refinement: *MSC/AFC Diffractometer Control Software*; data reduction: *CrystalStructure* (Molecular Structure Corporation & Rigaku, 2001); program(s) used

to solve structure: *SIR92* (Altomare *et al.*, 1994); program(s) used to refine structure: *CRYSTALS* (Betteridge *et al.*, 2003); molecular graphics: *ORTEP3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *CrystalStructure*.

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