C2-S1-C5	91.9(1)	C8a—S7a—C6a	91.0 (3)
C3-C2-S1	110.6 (2)	C9aC8aS7a	115.0 (4)
C2 ⁱ -C2-S1	120.9 (2)	C10a-C6a-S7a	107.7 (6)
C5-C4-C41	124.7 (2)	C5C6aS7a	120.8 (4)
C3-C4-C41	124.2 (2)	C8b—S7b—C6b	87.2 (7)
C4C5S1	111.8 (2)	C9b—C8b—S7b	113.3 (12)
C6aC5S1	115.5 (3)	C10bC6bS7b	113.0 (9)
C6b-C5-S1	124.9 (5)	C5C6bS7b	108.8 (7)
c	4-C41-C42-C43	-176.6 (2)	
C	41C42C43C44	178.7 (2)	
C	42C43C44C45	-176.5 (2)	
C	43C44C45C46	176.6 (3)	
C	44—C45—C46—C47	-177.1 (3)	
C	45C46C47C48a	-177.0 (3)	
C	45C46C47C48 <i>b</i>	158.0 (6)	
C	46—C47—C48a—C49a	-177.6 (4)	
C	47-C48a-C49a-C50a	147.7 (6)	
C	46—C47—C48 <i>b</i> —C49 <i>b</i>	-171.8(7)	
C	47C48bC49bC50b	-128.6 (9)	
Symmetry co	ode: (i) $\frac{1}{2} - x, \frac{1}{2} - y, -$	z.	

The disorder of the thiophene ring and the decyl group were

treated separately. First, the major peaks in the ΔF map were found around the outer thiophene ring after the first cycles of refinement. Two models were built for the outer thiophene ring and restraints (same 1,2 and 1,3 distances as in the inner thiophene with s.u.) were employed in the subsequent refinements. A free variable 2 was given to the occupation factor of one of the two models. The subsequent refinements gave the proportion 65.0(4): 35.0(4) for the two models. For the decyl chain, two reasonable models for the last three C atoms were found from the ΔF map; a free variable was then given to the occupation factor of one of the two models. During the refinement, restraints of the 1.2 and 1.3 distances among atoms C47-C50 were applied. Suprisingly, a very similar proportion, 65:35, was obtained for the two models after refinement. Therefore, a common free variable was given to the two parts of disorder in the next refinement. The proportion was found to be 65:35 with an s.u. of 0.4. In the final refinement, the occupation factors were fixed at 0.65 and 0.35. All non-H atoms were refined anisotropically in the major conformer and isotropically, except for the S atoms, in the minor conformer. The same displacement parameter was applied to the three disordered atoms (C48-C50). H atoms were constrained to the parent site using a riding model and refined isotropically (C-H 0.93-0.98 Å). The isotropic displacement parameters were adjusted to $1.5U_{eq}$ for methyl C atoms or $1.2U_{eq}$ for non-methyl C atoms. A final verification of possible voids was performed using PLATON (Spek, 1995).

Data collection: CAD-4 Software (Enraf-Nonius, 1989). Cell refinement: CAD-4 Software. Data reduction: NRC-2 and NRC-2A (Ahmed et al., 1973). Program(s) used to solve structure: SHELXS96 (Sheldrick, 1990). Program(s) used to refine structure: NRCVAX (Gabe et al., 1989) and SHELXL96 (Sheldrick, 1996). Molecular graphics: ORTEPII (Johnson, 1976) in NRCVAX. Software used to prepare material for publication: NRCVAX and SHELXL96.

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trans-2,3-Diiodo-2-butene-1,4-diol

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Abstract

The crystal structure of *trans*-2,3-diiodo-2-butene-1,4diol, C₄H₆I₂O₂, has been refined in space group $I4_1/a$, using data collected at 150 K. Bond angles around the sp^2 -hybridized C atom show significant deviations from 120°.

Comment

The title compound, (I), was prepared by iodine addition to the corresponding acetylene as described by Iserson & Smith (1967). Addition of iodine to acetylenes results in *trans* isomers; however, *cis* products have also been reported (Hollins & Campos, 1979; Moreau & Commeyras, 1985). NMR studies of the stereochemical aspects of (I) were not successful (Priebe, 1997); therefore, a crystal structure analysis of (I) was undertaken.



There are no unusual bond lengths in this molecule. However, two of the three bond angles around the C1 atom show significant deviations from 120° , *i.e.* C1ⁱ—C1—C2 = 127.8 (1)° and I—C1—C2 = 112.1 (1)° [symmetry code: (i) $\frac{3}{2} - x$, $\frac{3}{2} - y$, $-\frac{1}{2} - z$]. A search in the Cambridge Structural Database (Allen

A search in the Cambridge Structural Database (Allen & Kennard, 1993) for diiodated double/aromatic bonds (no disorder, $R \leq 0.10$, atom coordinates reported) revealed a total of 15 structures, of which only two have an iodinated non-cyclic double bond, namely, tetraiodo-ethylene (Haywood & Shirley, 1977) and a pyrazine-tetraiodoethylene complex (Dahl & Hassel, 1968). The present structure thus presents the first diiodinated non-cyclic double bond.



Fig. 1. Molecular packing diagram, with the atom numbering, of the *trans*-2,3-diiodo-2-butene-1,4-diol structure viewed along the c axis. Hydrogen bonds and charge-transfer interactions are illustrated using broken lines and dots, respectively. Displacement ellipsoids are drawn at the 75% probability level and H atoms are arbitrarily scaled.

There is only one type of hydrogen bond in the crystal, resulting in an infinite $\cdots O - H \cdots O - H \cdots$ chain, taking advantage of the cooperativity effect (Jeffrey & Saenger, 1991). Due to the fourfold screw axis, this chain constitutes a hydrophilic column with a hydrogen-bonded helix running parallel to the short crystallographic c axis (Fig. 1). The $O \cdots H$ distance is 1.87 Å and the $O - H \cdots O$ angle is 172° . The O atom is also engaged in a relatively weak charge-transfer interaction: $O \cdots I$ 3.429 (1) Å (the sum of the van der Waals radii is 3.55 Å).

The voluminous I atoms form a separate column along the c axis. The molecular packing arrangement is illustrated in Fig. 1.

Experimental

Due to the instability of *trans*-2,3-diiodo-2-butene-1,4-diol in dimethyl sulfoxide when exposed to light, the crystallization was performed at room temperature in complete darkness. Crystals were grown by overnight diffusion of methanol into a saturated solution of *trans*-2,3-diiodo-2-butene-1,4-diol in dimethyl sulfoxide.

Crystal data

 $C_4H_6I_2O_2$ Mo $K\alpha$ radiation $M_r = 339.89$ $\lambda = 0.71073 \text{ Å}$ Tetragonal Cell parameters from 8192 $I4_1/a$ (second setting) reflections a = 13.727(1) Å $\mu = 8.133 \text{ mm}^{-1}$ c = 8.130(1) Å T = 150(2) K $V = 1531.9(1) \text{ Å}^3$ Block $0.30 \times 0.30 \times 0.10 \text{ mm}$ Z = 8Colourless $D_x = 2.947 \text{ Mg m}^{-3}$ D_m not measured

17 735 measured reflections

2891 independent reflections

2646 reflections with

 $I > 2\sigma(I)$

 $k = -27 \rightarrow 27$

 $l = -12 \rightarrow 16$

 $R_{int} = 0.023$ $\theta_{max} = 44.93^{\circ}$ $h = -27 \rightarrow 19$

Data collection

Siemens SMART CCD areadetector diffractometer ω scan Absorption correction: analytical *XPREP* (Siemens, 1996), followed by an empirical correction of intensities due to absorption by the glass fibre and glue (*SADABS*; Sheldrick, 1996) $T_{min} = 0.140, T_{max} = 0.428$

Refinement

Refinement on F^2 $\Delta \rho_{max} = 1.103$ e Å $^{-3}$ $R[F^2 > 2\sigma(F^2)] = 0.027$ $\Delta \rho_{min} = -0.711$ e Å $^{-3}$ $wR(F^2) = 0.053$ Extinction correction:S = 2.239SHELXTL (Sheldrick,2891 reflections1994)39 parametersExtinction coefficient:H atoms constrained0.0049 (2)

$w = 1/[\sigma^2(F_o^2) + (0.0086P)^2]$	Scattering factors from
where $P = (F_o^2 + 2F_c^2)/3$	International Tables for
$(\Delta/\sigma)_{\rm max} = 0.001$	Crystallography (Vol. C)

Table 1. Fractional atomic coordinates and equivalent isotropic displacement parameters (Å²)

$$U_{\text{eq}} = (1/3) \sum_i \sum_j U^{ij} a_i^* a_i^* \mathbf{a}_i \cdot \mathbf{a}_j$$

	х	у	z	U_{eq}
I 1	0.654070 (8)	0.687482 (7)	-0.006829(11)	0.02098 (3)
01	0.75744 (8)	0.90831 (7)	-0.00683 (12)	0.01740 (14)
Cl	0.75028 (8)	0.76063 (8)	-0.16952 (15)	0.01362 (15)
C2	0.81343 (9)	0.83352 (10)	-0.0813 (2)	0.0171 (2)

Table 2. Selected geon	etric parameters (A, °))
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II—C1	2.122 (1)	C1—C1 ⁱ	1.341 (2)	
01—C2	1.418 (2)	C1—C2	1.505 (2)	
C1 ⁱ —C1—C2	127.8 (1)	C2—C1—I1	112.1 (1)	
C1 ⁱ —C1—I1	120.1 (1)	O1—C2—C1	111.9 (1)	
C1 ⁱ —C1—C2—O1	117.9 (2)	I1—C1—C2—O1	-62.2(1)	
Symmetry code: (i) $\frac{3}{2} - x$, $\frac{3}{2} - y$, $-\frac{1}{2} - z$.				

The asymmetric unit contains half a molecule. All atoms, except for the three H atoms, were refined anisotropically. The two ethylene H atoms were fixed in idealized positions, refining only the C—H distance with $U_{iso}(H) = 1.2U_{eq}(C2)$. The hydroxy H atom was localized using the *AFIX* card 147 of *SHELXTL* (Sheldrick, 1994) and U_{iso} was set to $1.2U_{eq}(O1)$ in the subsequent refinement.

Data collection: *SMART* (Siemens, 1995). Cell refinement: *SAINT* (Siemens, 1995). Data reduction: *SAINT*. Program(s) used to solve structure: *SHELXTL*. Program(s) used to refine structure: *SHELXTL*.

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9-(4-Dimethylaminophenyl)-3,3,6,6-tetramethyl-3,4,6,7,9,10-hexahydro-1,8(2H,5H)acridinedione

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Abstract

In the title compound, $C_{25}H_{32}N_2O_2$, the central ring in the acridine moiety is nearly planar, while the two outer rings adopt half-chair conformations. The phenyl ring is axial with respect to the central pyridine ring.

Comment

Aminoacridinyl derivatives are biologically useful molecules, especially as potential anticancer agents. Acridines have been used as antibacterial agents for wound therapy (Neidle, 1979). Some of them fluoresce well as laser dyes in alcohol solvents (Selladurai *et al.*, 1990).

The bond lengths and angles of the title compound, (1), are similar to those of related acridine derivatives (Selladurai *et al.*, 1989, 1990; Sivaraman *et*



al., 1994, 1996). The O1 atom deviates significantly [-0.256(2) Å] from the mean plane of ring I (Fig. 1), while the deviation of the O2 atom [-0.036(5) Å] from the mean plane of ring III is small but significant. Phenyl ring IV is axial with respect to central pyridine ring II. The bonding geometry about each of the two N atoms is planar. Phenyl ring IV is nearly perpendicular to the mean plane of the acridine moiety. The buckling of the