

Butanedithiol diacetate

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Key indicators

Single-crystal X-ray study
 $T = 187\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$
Disorder in main residue
 R factor = 0.048
 wR factor = 0.134
Data-to-parameter ratio = 18.1

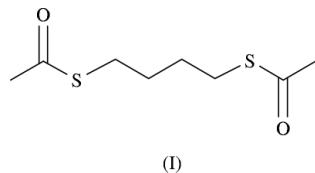
For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

In the solid state, molecules of butane-1,4-diyl bis(S-acetate), $\text{C}_8\text{H}_{14}\text{O}_2\text{S}_2$, are centrosymmetric and are associated *via* weak hydrogen bonds. The molecular structure, determined by X-ray crystallography, is in good agreement with that obtained by density functional geometry optimization.

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Comment

Thioesters are very important acetylating agents in biochemical processes, as well as in many chemical transformations (Nicolaou, 1977; Hirama *et al.*, 1979; Zheng *et al.*, 1999). We obtained butanedithiol diacetate, $\text{H}_3\text{CC(O)SCH}_2\text{CH}_2\text{CH}_2\text{CH}_2\text{SC(O)CH}_3$, (I), as a by-product in the synthesis of 1,4-butanedithiol monoacetate, $\text{HSCH}_2\text{CH}_2\text{CH}_2\text{CH}_2\text{SC(O)CH}_3$, (II), in a procedure similar to the preparation of ethanedithiol monoacetate and ethanedithiol diacetate (Wiesler *et al.*, 1996; Fleischer & Schollmeyer, 2001).



The molecular structure of (I), as found in the solid state, is depicted in Fig. 1; the molecule is centrosymmetric. Table 1 shows a comparison of selected structural parameters from the crystal structure with those obtained by a density functional (DF) geometry optimization. The molecular structure was first optimized at the Hartree–Fock level with a 6-31G(d) basis set, starting from the molecular structure as found in the solid state. Subsequently the structure was re-optimized employing density functional theory (DFT) and a larger basis set [B3LYP/6-311+G(2d,p)]. Apart from the two S–C and the C1–C1ⁱ distances [symmetry code: (i) $1-x, 1-y, 1-z$], the DF bond lengths and angles agree quite well with the experimental values. The torsion angles C1ⁱ–C1–C2–S3 and C1–C2–S3–C4 differ substantially between the solid state and the DF-calculated isolated molecule. We attribute the differences in the above-mentioned bond distances to the chosen level of theory, but the difference in torsion angles to intermolecular hydrogen bonds in the solid state [$\text{O}5^{\text{ii}} \cdots \text{C}6 = 3.497(3)\text{ \AA}$ and $\text{O}5^{\text{ii}} \cdots \text{H}–\text{C}6 = 168.1(2)^{\circ}$; symmetry code: (ii) $x, \frac{1}{2}-y, \frac{1}{2}+z$]. The structural parameters of (I) agree well with those found for other compounds containing an S-acetyl moiety (Fleischer & Schollmeyer, 2001; Evans *et al.*, 1999; Divjaković *et al.*, 1992; Mackay *et al.*, 1992; Huber *et al.*, 1984; Mattes & Waldmann, 1983; Mattes *et al.*, 1977; Kiel *et al.*,

1974). This implies that the *S*-acetyl fragment is a relatively rigid structural unit, a hypothesis supported by an analysis of the bonding situation in terms of natural bond orbitals (Fleischer & Schollmeyer, 2001).

Experimental

Butanedithiol diacetate, (I), was obtained as a by-product in the synthesis of 1,4-butanedithiol monoacetate, (II), in a procedure similar to the preparation of ethanedithiol monoacetate and ethanedithiol diacetate (Wiesler *et al.*, 1996; Fleischer & Schollmeyer, 2001). Crystals of (I) precipitated at 278 K from its solution in (II). They were washed with cold petroleum ether and identified by elemental analysis and ^1H NMR.

Crystal data

$\text{C}_8\text{H}_{14}\text{O}_2\text{S}_2$
 $M_r = 206.31$
Monoclinic, $P2_1/c$
 $a = 6.1998 (7)$ Å
 $b = 8.0292 (7)$ Å
 $c = 10.5650 (10)$ Å
 $\beta = 95.942 (4)$ °
 $V = 523.09 (9)$ Å 3
 $Z = 2$

$D_x = 1.310 \text{ Mg m}^{-3}$
Cu $\text{K}\alpha$ radiation
Cell parameters from 25 reflections
 $\theta = 57\text{--}73$ °
 $\mu = 4.31 \text{ mm}^{-1}$
 $T = 187 (2)$ K
Block, colorless
 $0.25 \times 0.20 \times 0.15$ mm

Data collection

Enraf–Nonius CAD-4 diffractometer
 $\omega/2\theta$ scans
Absorption correction: ψ scan (*CORINC*; Dräger & Gattow, 1971)
 $T_{\min} = 0.383$, $T_{\max} = 0.524$
1126 measured reflections
1067 independent reflections

908 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.169$
 $\theta_{\text{max}} = 73.8$ °
 $h = -7 \rightarrow 7$
 $k = 0 \rightarrow 10$
 $l = 0 \rightarrow 13$
3 standard reflections frequency: 60 min intensity decay: 5%

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.048$
 $wR(F^2) = 0.134$
 $S = 1.12$
1067 reflections
59 parameters

H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0915P)^2]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.34 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.84 \text{ e } \text{\AA}^{-3}$

Table 1

Comparison of selected structural parameters (Å, °) from the molecular structure of (I) in the solid state (XRD) and from density functional (DF) geometry optimization [$\text{B3LYP}/6-311+\text{G}(2d,p)$] (Frisch *et al.*, 1995).

	XRD	DF
C1—Cl ¹	1.517 (4)	1.534
C1—C2	1.523 (3)	1.527
C2—S3	1.809 (2)	1.834
S3—C4	1.766 (3)	1.795
C4—C6	1.506 (3)	1.512
C4—O5	1.207 (3)	1.205
Cl ¹ —C1—C2	113.3 (2)	111.6
C1—C2—S3	113.6 (2)	110.4
C2—S3—C4	101.0 (1)	100.1
S3—C4—C6	113.2 (2)	113.6
O5—C4—C6	123.7 (3)	123.5
O5—C4—S3	123.1 (2)	122.9
Cl ¹ —C1—C2—S3	67.6 (3)	179.9
C1—C2—S3—C4	82.7 (2)	182.1
C2—S3—C4—O5	0.8 (2)	0.3
C2—S3—C4—C6	179.5 (2)	180.5

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

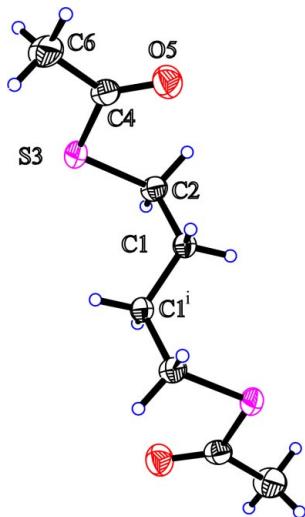


Figure 1

ORTEPII (Johnson, 1976) view of the centrosymmetric molecule of (I). Displacement ellipsoids are drawn at the 50% probability level. H atoms are shown as spheres of fixed radius. The methyl group is disordered (0.52:0.48, ratio fixed) and only one set of H atoms bound to C6 is shown.

Chemically equivalent H atoms were placed at calculated positions and were refined isotropically with the same displacement parameters using riding motion. The two different groups of H atoms resulting from the disorder of the methyl group were treated with two independent displacement parameters.

Data collection: *CAD-4 Software* (Enraf–Nonius, 1989); cell refinement: *CAD-4 Software*; data reduction: *CORINC* (Dräger & Gattow, 1971); program(s) used to solve structure: *SIR92* (Altomare *et al.*, 1994); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLATON* (Spek, 2001); software used to prepare material for publication: *SHELXL97*.

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