

1-O-Benzyl-2,3-O-isopropylidene-6-O-tosyl- α -L-sorbofuranose

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Received 9 May 2013; accepted 5 June 2013

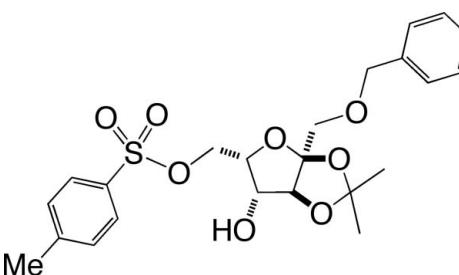
Key indicators: single-crystal X-ray study; $T = 150\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$; R factor = 0.044; wR factor = 0.127; data-to-parameter ratio = 15.9.

In the title compound (systematic name: {(3aS,5S,6R,6aS)-3a-[(benzyloxy)methyl]-6-hydroxy-2,2-dimethyltetrahydrofuro-[2,3-*d*][1,3]dioxol-5-yl)methyl 4-methylbenzenesulfonate}, $C_{23}H_{28}O_8S$, the absolute structure and relative stereochemistry of the four chiral centres have been established by X-ray crystallography, with the absolute configuration inferred from the use of L-sorbose as the starting material. The central furanose ring adopts a slightly twisted envelope conformation (with the C atom bearing the methylbenzenesulfonate substituent as the flap) from which three substituents depart pseudo-axially ($-\text{CH}_2-\text{O}-\text{benzyl}$, $-\text{OH}$ and one acetonide O atom) and two substituents pseudo-equatorially ($-\text{CH}_2-\text{O}-\text{tosyl}$ and second acetonide O atom). The dioxalane ring is in a flattened envelope conformation with the fused CH C atom as the flap. In the crystal, molecules pack in columns along [010] linked by $\text{O}-\text{H}\cdots\text{O}$ hydrogen bonds involving the furanose hydroxy group and furanose ether O atom.

Related literature

The title compound is a novel intermediate in the synthesis of 1-deoxynojirimycin (DNJ) analogues. For examples of the use of monosaccharide starting materials in iminosugar syntheses, see: Compain & Martin (2001); Cipolla *et al.* (2003); Best, Wang *et al.* (2010); Wilkinson *et al.* (2010); Nash *et al.* (2011); Zhang *et al.* (2011); Lenagh-Snow *et al.* (2011); Simone *et al.* (2012); Soengas *et al.* (2012); Kato *et al.* (2012). For examples of the synthesis of other biologically active compounds from monosaccharides, see: Compain *et al.* (2009); Sridhar *et al.* (2012); Das *et al.* (2012); Dhavale & Matin (2005); Compain & Martin (2001); Derosa & Maffioli (2012); Lew *et al.* (2000); Itzstein *et al.* (1993). For glycosidase inhibitors, see: Houston & Blanchfield (2003). For iminosugars as glycosidase inhibitors, see: Zechel *et al.* (2003); de Melo *et al.* (2006); Compain &

Martin (2007). For examples of the clinical uses of iminosugars, see: Cox *et al.* (2003); Venier *et al.* (2012); Derosa & Maffioli (2012). For iminosugars in the treatment of cancer, cystic fibrosis and viral diseases, see: Nishimura (2003); Lawton & Witty (2011); Best, Jenkinson *et al.* (2010); Compain & Martin (2007); Pollock *et al.* (2008). For the syntheses of DNJ and its analogues from L-sorbose, see: Beaupere *et al.* (1989); Masson *et al.* (2000); Tamayo *et al.* (2010); O'Brien & Murphy (2011). For the synthesis of 1-O-benzoyl-2,3-O-isopropylidene-6-O-tosyl- α -L-sorbofuranose, which bears structural similarity to the title compound, see: Fehér & Varga (1966).



Experimental

Crystal data

$C_{23}H_{28}O_8S$	$V = 2321.04(6)\text{ \AA}^3$
$M_r = 464.51$	$Z = 4$
Monoclinic, $C2$	$Cu K\alpha$ radiation
$a = 22.6192(3)\text{ \AA}$	$\mu = 1.64\text{ mm}^{-1}$
$b = 5.5649(1)\text{ \AA}$	$T = 150\text{ K}$
$c = 19.0631(3)\text{ \AA}$	$0.29 \times 0.06 \times 0.02\text{ mm}$
$\beta = 104.696(2)$	

Data collection

Agilent SuperNova (Dual, Cu at zero, Atlas) diffractometer	24544 measured reflections
Absorption correction: multi-scan (<i>CrysAlis PRO</i> ; Agilent, 2011)	4672 independent reflections
$T_{\min} = 0.628$, $T_{\max} = 1.000$	4541 reflections with $I > 2\sigma(I)$
	$R_{\text{int}} = 0.031$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.044$	H-atom parameters constrained
$wR(F^2) = 0.127$	$\Delta\rho_{\text{max}} = 0.48\text{ e \AA}^{-3}$
$S = 1.16$	$\Delta\rho_{\text{min}} = -0.27\text{ e \AA}^{-3}$
4672 reflections	Absolute structure: Flack (1983), 2165 Friedel pairs
293 parameters	Flack parameter: 0.000 (15)
1 restraint	

Table 1
Hydrogen-bond geometry (\AA , $^\circ$).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
O3—H3O \cdots O4 ⁱ	0.84	1.98	2.812 (2)	174

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

Data collection: *CrysAlis PRO* (Agilent, 2011); cell refinement: *CrysAlis PRO*; data reduction: *CrysAlis PRO*; program(s) used to solve structure: *SIR97* (Altomare *et al.*, 1999); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *Xtal3.6* (Hall *et al.*, 1999), *ORTEPII* (Johnson, 1976), *SHELXLE* (Hübschle *et al.*, 2011), *Mercury* (Macrae *et al.*, 2006) and *WinGX*

(Farrugia, 2012); software used to prepare material for publication: *publCIF* (Westrip, (2010).

The University of Sydney is gratefully acknowledged for funding.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: LH5615).

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supplementary materials

Acta Cryst. (2013). E69, o1069–o1070 [doi:10.1107/S1600536813015638]

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Comment

Monosaccharides provide a vast and formidable chiral pool of starting materials, whose utilization continues to expand in the enantiospecific syntheses of natural products (Sridhar *et al.*, 2012; Das *et al.*, 2012), in particular of carbohydrate mimetics of the carbasugar (Derosa & Maffioli, 2012; Lew *et al.* 2000), C-glycoside (Compain & Martin, 2001; Dhavale & Matin, 2005; Compain *et al.*, 2009), THP (Itzstein *et al.* 1993) and iminosugar (Cipolla *et al.*, 2003; Wilkinson *et al.* (2010); Nash *et al.*, 2011; Zhang *et al.*, 2011; Lenagh-Snow *et al.*, 2011; Simone *et al.*, 2012; Soengas *et al.* 2012; Best, Wang *et al.* 2010; Kato *et al.*, 2012) types, to mention a few.

Iminosugars have been recognized as a class of potent inhibitors of glycosidase enzymes (Houston & Blanchfield 2003; Zechel *et al.*, 2003; de Melo *et al.*, 2006; Compain and Martin, 2007). These potent biological activities have culminated in the marketing of *N*-butyl-DNJ for the treatment of Gauchers disease (Miglustat) (Cox *et al.*, 2003; Venier *et al.*, 2012), *N*-hydroxyethyl-DNJ for type II diabetes (Miglitol) (Derosa & Maffioli, 2012) while other iminosugars have opened up new in-roads in the treatment of cancer (Nishimura, 2003; Lawton & Witty 2011), cystic fibrosis (Best, Jenkinson *et al.*, 2010) and as antivirals (Compain & Martin, 2007; Pollock *et al.* 2008).

The first synthesis of DNJ (3 in Fig. 1) from starting material L-sorbose (1) utilized triphenylphosphine, carbon tetrabromide and lithium azide to effect the key transformation which installs an azido group in place of the C5 hydroxy (Beaupere *et al.*, 1989). Syntheses of further DNJ derivatives from L-sorbose have been reported (Masson *et al.*, 2000; Tamayo *et al.*, 2010; O'Brien & Murphy, 2011). The title compound (2 in Fig. 1) bears orthogonal protecting groups on four of the five hydroxy groups, thus opening up points of synthetic divergence to novel classes of iminosugar glycomimetics based on a DNJ scaffold.

In the title compound (Fig. 2), the central furanose ring adopts a slightly twisted envelope conformation with C4 forming the flap. The O1 and C5 substituents are positioned pseudo-equatorially, while the C1', O2 and O3 substituents are positioned pseudo-axially. The dioxalane ring is in a flattened envelope conformation with C2 forming the flap. The title compound bears structural similarity to 1-*O*-benzoyl-2,3-*O*-isopropylidene-6-*O*-tosyl- α -L-sorbofuranose ($[\alpha]_D^{20} 0^\circ$ ($c = 1$ in CHCl_3); m.p. 428–429 K (decomposition)) (Fehér & Vargha, 1966). In the crystal, molecules pack in columns in the [010] direction linked by O—H···O hydrogen bonds involving the furanose hydroxy group and furanose ether oxygen atom (Fig. 3).

Experimental

Triethylamine (3.43 ml, 24.6 mmol), 4-dimethylaminopyridine (120 mg, 982 μmol), and 4-toluenesulfonyl chloride (2.35 g, 12.3 mmol) were carefully added in succession to a stirring solution of 1-*O*-benzyl-2,3-*O*-isopropylidene- α -L-sorbofuranose (3.06 g, 9.84 mmol) in dichloromethane (200 ml) under an inert atmosphere. The reaction mixture was stirred for 28 h at room temperature, after which, analysis by TLC (acetone/hexane 1:2) showed total consumption of the starting material ($R_f = 0.18$) and formation of a UV-active product ($R_f = 0.44$). The crude mixture was washed with an aqueous

hydrochloric acid solution (1 M , 100 ml) and the organic layers collected, dried over magnesium sulfate, filtered and concentrated *in vacuo*. The compound crystallized on standing as a pale yellow solid in quantitative yield (4.57 g).

Refinement

All H atoms attached to C atoms were positioned geometrically, and allowed to ride on their parent atoms, with C—H bond lengths of 0.95 Å (Ar—H), 1.0 (CH), 0.99 Å (CH₂) or 0.98 Å (CH₃), and isotropic displacement parameters set to 1.2 (CH and CH₂) or 1.5 times (CH₃) the U_{eq} of the parent atom. The H atom attached to O₃ was positioned geometrically and allowed to ride on the parent atom, with an O—H bond length of 0.84 Å and isotropic displacement parameters set to 1.5 $U_{\text{eq}}(\text{O})$.

Computing details

Data collection: *CrysAlis PRO* (Agilent, 2011); cell refinement: *CrysAlis PRO* (Agilent, 2011); data reduction: *CrysAlis PRO* (Agilent, 2011); program(s) used to solve structure: *SIR97* (Altomare *et al.*, 1999); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *Xtal3.6* (Hall *et al.*, 1999), *ORTEPII* (Johnson, 1976), *SHELXLE* (Hübschle *et al.*, 2011), Mercury (Macrae *et al.*, 2006) and *WinGX* (Farrugia, 2012); software used to prepare material for publication: *publCIF* (Westrip, (2010)).

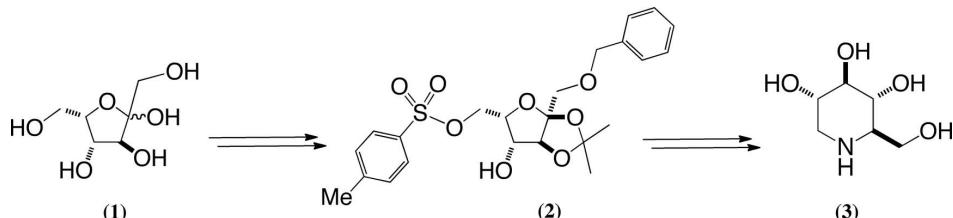


Figure 1

Relationship of L-sorbose (1) starting material to title compound (2) and deoxynojirimycin (DNJ) (3)

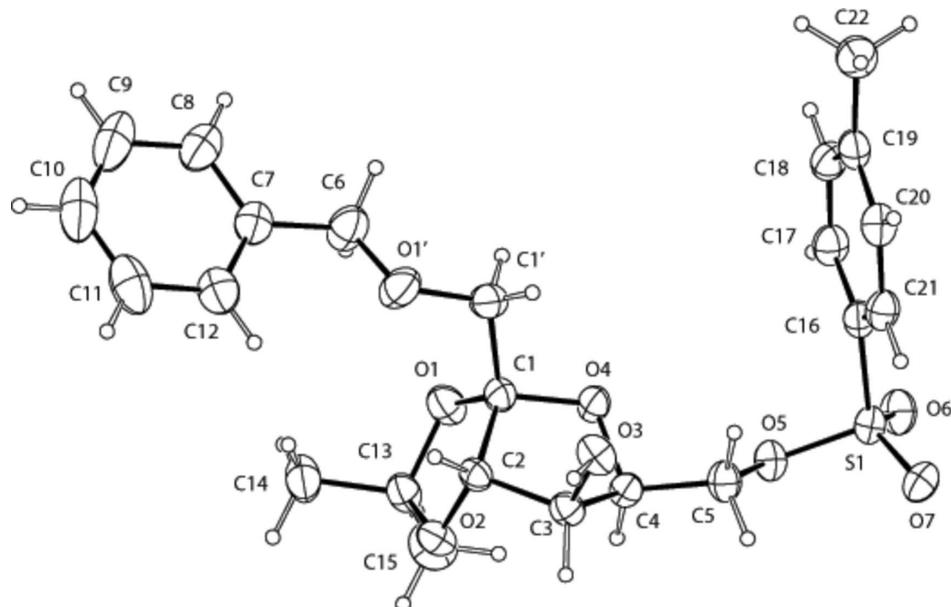
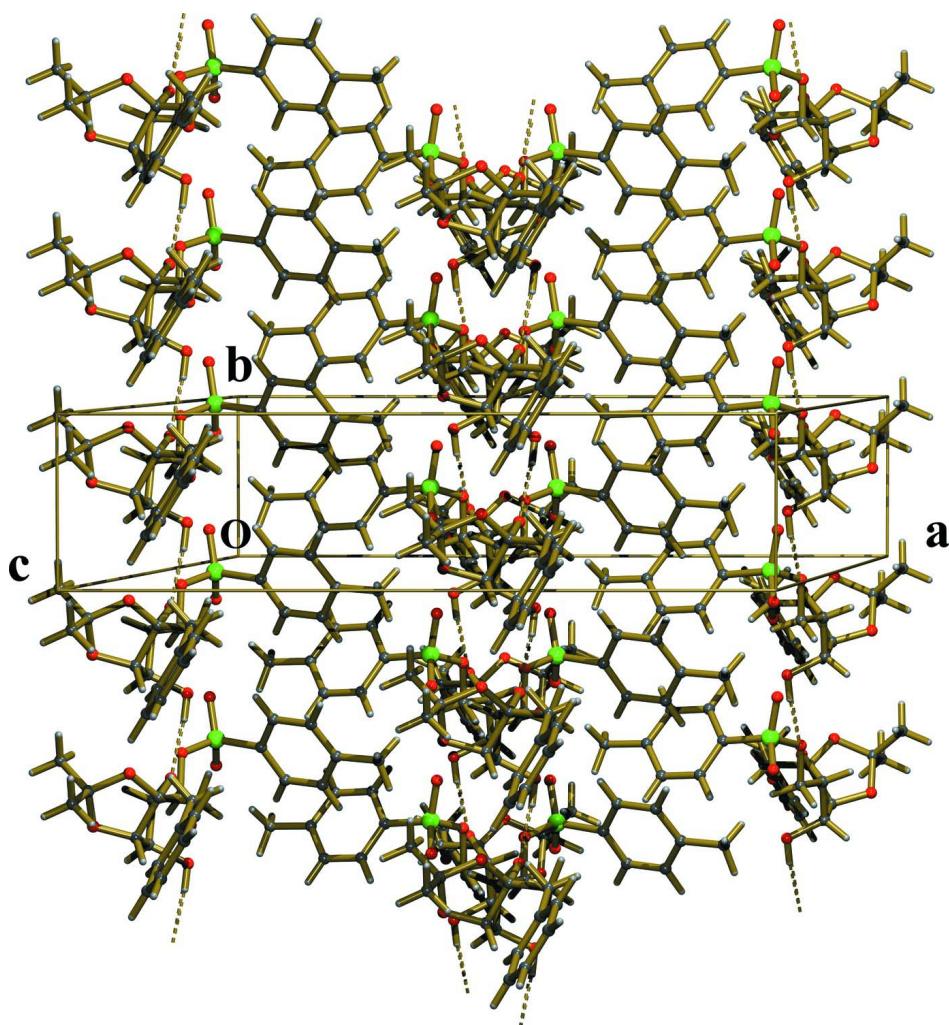


Figure 2

The molecular structure of the title compound, with anisotropic displacement ellipsoids shown at the 50% probability level.

**Figure 3**

Molecular packing diagram (Macrae *et al.*, 2006) with respect to the unit cell. The molecules stack in columns parallel to the *b* axis, with molecules within a column linked by intermolecular hydrogen bonds (dashed lines) between the hydroxy O3 moiety and the furanose oxygen O4.

{(3a*S*,5*S*,6*R*,6a*S*)-3a-[(Benzyl oxy)methyl]-6-hydroxy-2,2-dimethyltetrahydrofuro[2,3-*d*][1,3]dioxol-5-yl}methyl 4-methylbenzenesulfonate

Crystal data

C₂₃H₂₈O₈S
 $M_r = 464.51$
 Monoclinic, C2
 Hall symbol: C 2y
 $a = 22.6192 (3)$ Å
 $b = 5.5649 (1)$ Å
 $c = 19.0631 (3)$ Å
 $\beta = 104.696 (2)^\circ$
 $V = 2321.04 (6)$ Å³
 $Z = 4$

$F(000) = 984$
 $D_x = 1.329 \text{ Mg m}^{-3}$
 Cu $K\alpha$ radiation, $\lambda = 1.5418$ Å
 Cell parameters from 13737 reflections
 $\theta = 4.0\text{--}76.2^\circ$
 $\mu = 1.64 \text{ mm}^{-1}$
 $T = 150 \text{ K}$
 Blade, colourless
 $0.29 \times 0.06 \times 0.02$ mm

Data collection

Agilent SuperNova (Dual, Cu at zero, Atlas) diffractometer
 Radiation source: SuperNova (Cu) X-ray Source
 Mirror monochromator
 Detector resolution: 10.5861 pixels mm⁻¹
 ω scans
 Absorption correction: multi-scan (*CrysAlis PRO*; Agilent, 2011)

$T_{\min} = 0.628$, $T_{\max} = 1.000$
 24544 measured reflections
 4672 independent reflections
 4541 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.031$
 $\theta_{\max} = 76.4^\circ$, $\theta_{\min} = 4.0^\circ$
 $h = -28 \rightarrow 28$
 $k = -7 \rightarrow 6$
 $l = -24 \rightarrow 24$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.044$
 $wR(F^2) = 0.127$
 $S = 1.16$
 4672 reflections
 293 parameters
 1 restraint
 Primary atom site location: structure-invariant direct methods
 Secondary atom site location: difference Fourier map

Hydrogen site location: inferred from neighbouring sites
 H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.1P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.48 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.27 \text{ e } \text{\AA}^{-3}$
 Absolute structure: Flack (1983), 2165 Friedel pairs
 Flack parameter: 0.000 (15)

Special details

Experimental. Analysis: $[\alpha]_D^{25} 0.20^\circ$ (*c* 0.2 in CHCl_3); IR (KBr, cm^{-1}): 3594-3205 (*br*, OH), 3095, 3046 (w, ArC-H), 2984, 2950, 2925, 2864 (st, alkyl C-H), 1369, 1178 (s, S=O); ¹H NMR (CDCl_3 , 400 MHz, p.p.m.): 1.27, 1.46 [2 x 3H, 2 x s, $\text{C}(\text{CH}_3)_2$], 2.43 [3H, s, TsCH_3], 3.49 [1H, d, $J_{\text{H}_3,\text{H}_4} = 11.8 \text{ Hz}$, H_3], 3.57 [1H, d, $J_{\text{H}_1,\text{H}_1'} = 10.1 \text{ Hz}$, H_1], 3.76 [1H, d, $J_{\text{H}_1',\text{H}_1} = 10.0 \text{ Hz}$, H_1'], 4.05 [1H, dd, $J_{\text{H}_4,\text{H}_3} = 11.5 \text{ Hz}$, H_4], 4.16 [1H, dd, $J_{\text{H}_6,\text{H}_5} = 6.9 \text{ Hz}$, H_6], 4.30 [1H, dd, $J_{\text{H}_6',\text{H}_5} = 4.9$, H_6'], 4.36 [1H, s, OH], 4.40 [1H, ddd, $J_{\text{H}_5,\text{H}_4} = 2.5 \text{ Hz}$, $J_{\text{H}_5,\text{H}_6} = 4.7 \text{ Hz}$, $J_{\text{H}_5,\text{H}_6} = 7.2 \text{ Hz}$, H_5], 4.52 [1H, d, $J_{\text{CHAHB},\text{CHAHB}} = 11.7 \text{ Hz}$, CH_AH_B (Bn)], 4.59 [1H, d, $J_{\text{CHAHB},\text{CHAHB}} = 11.7 \text{ Hz}$, CH_AH_B (Bn)], 7.22-7.26 [2H, m, 2 x ArHs (Bn-*o*)], 7.30-7.37 [5H, m, 2 x ArHs (Ts) and 3 x ArHs (Bn-*m,p*)], 7.81 [2H, d, ArHs (Ts)]; ¹³C NMR (CDCl_3 , 100 MHz, p.p.m.): 21.7 [CH_3 (Ts)], 26.1, 27.2 [2 x CH_3], 68.1 [C6], 71.3 [C1], 74.2 [CH_2 (Bn)], 74.4 [C4], 79.8 [C5], 86.4 [C3], 112.8 [C2], 112.9 [Cq acetonide], 128.0 [2 x ArCs (Bn-*o*)], 128.2 [2 x ArCs (Ts)], 128.5 [1 x ArC (Bn-*p*)], 128.8, 129.9 [2 x ArCs (Bn-*m*) and 2 x ArCs (Ts)], 133.0 [Cq- CH_3], 136.5 [Cq (Bn)], 144.9 [Cq-S]; HRMS (ESI⁺): found 487.13977 [$\text{M}+\text{Na}$]⁺ $\text{C}_{23}\text{H}_{28}\text{NaO}_8\text{S}$, requires 487.13971. M.p.: 377-378K

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted *R*-factor *wR* and goodness of fit *S* are based on F^2 , conventional *R*-factors *R* are based on *F*, with *F* set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating *R*-factors(gt) etc. and is not relevant to the choice of reflections for refinement. *R*-factors based on *F²* are statistically about twice as large as those based on *F*, and *R*-factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^* / U_{\text{eq}}$
S1	0.404262 (19)	0.50470 (8)	0.48560 (2)	0.02722 (14)
O1	0.44717 (7)	0.4027 (3)	0.17231 (8)	0.0355 (3)
O1'	0.34123 (8)	0.1296 (4)	0.09744 (8)	0.0438 (4)
O2	0.50981 (6)	0.0816 (3)	0.21233 (8)	0.0313 (3)
O3	0.39754 (6)	-0.1689 (3)	0.29897 (8)	0.0316 (3)

H3O	0.4000	-0.3170	0.2916	0.047*
O4	0.41584 (6)	0.3389 (2)	0.27843 (7)	0.0279 (3)
O5	0.43901 (6)	0.4404 (3)	0.42627 (7)	0.0289 (3)
O6	0.41714 (7)	0.7530 (3)	0.50128 (8)	0.0354 (3)
O7	0.41841 (7)	0.3306 (3)	0.54278 (7)	0.0345 (3)
C1	0.41496 (8)	0.2518 (4)	0.20750 (9)	0.0269 (4)
C1'	0.34827 (9)	0.2402 (5)	0.16634 (11)	0.0383 (5)
H1'A	0.3252	0.1477	0.1949	0.046*
H1'B	0.3312	0.4048	0.1596	0.046*
C2	0.45054 (8)	0.0140 (4)	0.21843 (9)	0.0259 (3)
H2	0.4311	-0.1123	0.1826	0.031*
C3	0.45313 (8)	-0.0559 (4)	0.29689 (10)	0.0270 (4)
H3	0.4895	-0.1575	0.3191	0.032*
C4	0.45641 (8)	0.1905 (3)	0.33235 (9)	0.0254 (4)
H4	0.4991	0.2542	0.3426	0.030*
C5	0.43459 (9)	0.1918 (4)	0.40058 (10)	0.0302 (4)
H5A	0.3918	0.1348	0.3904	0.036*
H5B	0.4605	0.0855	0.4376	0.036*
C6	0.32399 (13)	0.2890 (5)	0.03842 (12)	0.0436 (5)
H6A	0.3540	0.4216	0.0443	0.052*
H6B	0.2835	0.3594	0.0370	0.052*
C7	0.32113 (10)	0.1574 (5)	-0.03123 (11)	0.0365 (5)
C8	0.28553 (11)	0.2509 (6)	-0.09630 (12)	0.0459 (6)
H8	0.2630	0.3950	-0.0961	0.055*
C9	0.28292 (15)	0.1343 (7)	-0.16111 (13)	0.0602 (9)
H9	0.2589	0.1997	-0.2052	0.072*
C10	0.31489 (15)	-0.0766 (7)	-0.16223 (15)	0.0611 (9)
H10	0.3129	-0.1558	-0.2069	0.073*
C11	0.34979 (14)	-0.1721 (6)	-0.09822 (17)	0.0563 (7)
H11	0.3716	-0.3177	-0.0987	0.068*
C12	0.35291 (11)	-0.0536 (5)	-0.03250 (13)	0.0428 (5)
H12	0.3771	-0.1190	0.0115	0.051*
C13	0.50589 (9)	0.3003 (4)	0.17266 (11)	0.0313 (4)
C14	0.50694 (13)	0.2538 (6)	0.09465 (13)	0.0496 (6)
H14A	0.4735	0.1446	0.0721	0.074*
H14B	0.5461	0.1807	0.0934	0.074*
H14C	0.5019	0.4060	0.0679	0.074*
C15	0.55631 (12)	0.4642 (5)	0.21200 (15)	0.0481 (6)
H15A	0.5959	0.3946	0.2112	0.072*
H15B	0.5541	0.4834	0.2624	0.072*
H15C	0.5518	0.6215	0.1881	0.072*
C16	0.32657 (8)	0.4727 (4)	0.44051 (9)	0.0281 (4)
C17	0.29721 (10)	0.6518 (4)	0.39353 (11)	0.0324 (4)
H17	0.3195	0.7858	0.3828	0.039*
C18	0.23503 (10)	0.6319 (5)	0.36260 (11)	0.0355 (4)
H18	0.2147	0.7545	0.3308	0.043*
C19	0.20172 (9)	0.4365 (4)	0.37695 (10)	0.0329 (4)
C20	0.23236 (10)	0.2542 (4)	0.42171 (11)	0.0332 (4)
H20	0.2105	0.1161	0.4302	0.040*

C21	0.29487 (9)	0.2718 (4)	0.45432 (10)	0.0307 (4)
H21	0.3154	0.1482	0.4855	0.037*
C22	0.13348 (10)	0.4233 (6)	0.34609 (13)	0.0456 (6)
H22A	0.1234	0.4671	0.2946	0.068*
H22B	0.1133	0.5349	0.3723	0.068*
H22C	0.1194	0.2593	0.3513	0.068*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.0334 (2)	0.0263 (3)	0.0226 (2)	-0.00197 (16)	0.00836 (15)	-0.00147 (16)
O1	0.0404 (7)	0.0297 (8)	0.0412 (7)	0.0075 (6)	0.0190 (6)	0.0082 (6)
O1'	0.0522 (9)	0.0409 (10)	0.0313 (7)	0.0102 (7)	-0.0020 (6)	-0.0056 (7)
O2	0.0278 (6)	0.0284 (8)	0.0400 (7)	0.0039 (5)	0.0129 (5)	0.0037 (6)
O3	0.0356 (7)	0.0208 (7)	0.0417 (7)	-0.0025 (5)	0.0161 (6)	-0.0016 (5)
O4	0.0374 (7)	0.0213 (7)	0.0243 (6)	0.0056 (5)	0.0067 (5)	-0.0019 (5)
O5	0.0347 (6)	0.0267 (8)	0.0275 (6)	-0.0050 (5)	0.0121 (5)	-0.0029 (5)
O6	0.0441 (7)	0.0297 (8)	0.0337 (6)	-0.0060 (6)	0.0123 (6)	-0.0082 (6)
O7	0.0395 (7)	0.0384 (9)	0.0246 (6)	-0.0005 (6)	0.0061 (5)	0.0043 (6)
C1	0.0299 (8)	0.0266 (10)	0.0246 (8)	0.0038 (7)	0.0075 (7)	-0.0011 (7)
C1'	0.0326 (9)	0.0459 (13)	0.0328 (9)	0.0097 (9)	0.0017 (8)	-0.0080 (9)
C2	0.0265 (7)	0.0232 (9)	0.0287 (8)	0.0022 (7)	0.0083 (6)	-0.0027 (7)
C3	0.0288 (8)	0.0225 (10)	0.0305 (8)	0.0028 (7)	0.0090 (6)	0.0013 (7)
C4	0.0292 (8)	0.0217 (9)	0.0249 (8)	0.0005 (7)	0.0059 (6)	-0.0003 (7)
C5	0.0380 (9)	0.0256 (10)	0.0291 (8)	-0.0023 (7)	0.0125 (7)	-0.0020 (7)
C6	0.0591 (13)	0.0344 (13)	0.0342 (10)	0.0076 (11)	0.0061 (9)	0.0002 (9)
C7	0.0381 (10)	0.0383 (12)	0.0337 (10)	-0.0063 (9)	0.0102 (8)	-0.0017 (9)
C8	0.0475 (12)	0.0535 (16)	0.0346 (10)	-0.0093 (11)	0.0065 (9)	0.0032 (10)
C9	0.0682 (17)	0.081 (2)	0.0319 (11)	-0.0281 (17)	0.0130 (11)	-0.0017 (13)
C10	0.0691 (17)	0.079 (2)	0.0430 (12)	-0.0328 (17)	0.0275 (12)	-0.0210 (14)
C11	0.0620 (15)	0.0521 (16)	0.0651 (16)	-0.0171 (13)	0.0350 (13)	-0.0228 (14)
C12	0.0424 (11)	0.0429 (15)	0.0447 (11)	-0.0075 (10)	0.0143 (9)	-0.0080 (10)
C13	0.0350 (9)	0.0299 (11)	0.0327 (9)	0.0028 (8)	0.0153 (7)	0.0020 (8)
C14	0.0604 (14)	0.0579 (17)	0.0359 (11)	0.0145 (13)	0.0223 (10)	0.0010 (11)
C15	0.0476 (12)	0.0399 (15)	0.0595 (14)	-0.0129 (10)	0.0182 (10)	-0.0004 (11)
C16	0.0327 (8)	0.0286 (11)	0.0247 (7)	0.0014 (7)	0.0105 (6)	-0.0002 (7)
C17	0.0405 (10)	0.0279 (11)	0.0305 (8)	-0.0002 (8)	0.0122 (7)	0.0037 (8)
C18	0.0419 (10)	0.0355 (12)	0.0287 (9)	0.0067 (9)	0.0081 (7)	0.0033 (8)
C19	0.0347 (9)	0.0385 (12)	0.0273 (8)	0.0014 (8)	0.0113 (7)	-0.0059 (8)
C20	0.0391 (10)	0.0307 (11)	0.0319 (9)	-0.0055 (8)	0.0127 (7)	-0.0024 (8)
C21	0.0385 (9)	0.0284 (11)	0.0264 (8)	-0.0018 (8)	0.0104 (7)	0.0022 (7)
C22	0.0367 (10)	0.0585 (16)	0.0400 (11)	-0.0008 (10)	0.0069 (8)	-0.0096 (11)

Geometric parameters (\AA , $^\circ$)

S1—O6	1.4278 (17)	C8—C9	1.384 (4)
S1—O7	1.4326 (16)	C8—H8	0.9500
S1—O5	1.5736 (13)	C9—C10	1.382 (6)
S1—C16	1.7585 (19)	C9—H9	0.9500
O1—C1	1.391 (2)	C10—C11	1.380 (5)

O1—C13	1.444 (2)	C10—H10	0.9500
O1'—C6	1.408 (3)	C11—C12	1.402 (4)
O1'—C1'	1.422 (3)	C11—H11	0.9500
O2—C13	1.424 (3)	C12—H12	0.9500
O2—C2	1.425 (2)	C13—C15	1.504 (3)
O3—C3	1.415 (2)	C13—C14	1.516 (3)
O3—H3O	0.8400	C14—H14A	0.9800
O4—C1	1.432 (2)	C14—H14B	0.9800
O4—C4	1.450 (2)	C14—H14C	0.9800
O5—C5	1.463 (2)	C15—H15A	0.9800
C1—C1'	1.515 (3)	C15—H15B	0.9800
C1—C2	1.535 (3)	C15—H15C	0.9800
C1'—H1'A	0.9900	C16—C21	1.389 (3)
C1'—H1'B	0.9900	C16—C17	1.391 (3)
C2—C3	1.532 (2)	C17—C18	1.385 (3)
C2—H2	1.0000	C17—H17	0.9500
C3—C4	1.522 (3)	C18—C19	1.389 (3)
C3—H3	1.0000	C18—H18	0.9500
C4—C5	1.504 (2)	C19—C20	1.392 (3)
C4—H4	1.0000	C19—C22	1.508 (3)
C5—H5A	0.9900	C20—C21	1.396 (3)
C5—H5B	0.9900	C20—H20	0.9500
C6—C7	1.503 (3)	C21—H21	0.9500
C6—H6A	0.9900	C22—H22A	0.9800
C6—H6B	0.9900	C22—H22B	0.9800
C7—C12	1.380 (4)	C22—H22C	0.9800
C7—C8	1.397 (3)		
O6—S1—O7	120.05 (10)	C9—C8—C7	120.2 (3)
O6—S1—O5	104.95 (8)	C9—C8—H8	119.9
O7—S1—O5	109.71 (9)	C7—C8—H8	119.9
O6—S1—C16	109.01 (10)	C10—C9—C8	120.6 (3)
O7—S1—C16	107.82 (9)	C10—C9—H9	119.7
O5—S1—C16	104.19 (8)	C8—C9—H9	119.7
C1—O1—C13	110.70 (15)	C11—C10—C9	119.8 (3)
C6—O1'—C1'	114.1 (2)	C11—C10—H10	120.1
C13—O2—C2	109.61 (14)	C9—C10—H10	120.1
C3—O3—H3O	109.5	C10—C11—C12	119.8 (3)
C1—O4—C4	109.34 (14)	C10—C11—H11	120.1
C5—O5—S1	116.84 (11)	C12—C11—H11	120.1
O1—C1—O4	111.57 (16)	C7—C12—C11	120.5 (3)
O1—C1—C1'	110.53 (16)	C7—C12—H12	119.7
O4—C1—C1'	106.07 (14)	C11—C12—H12	119.7
O1—C1—C2	105.38 (14)	O2—C13—O1	105.86 (14)
O4—C1—C2	106.41 (14)	O2—C13—C15	108.44 (18)
C1'—C1—C2	116.89 (18)	O1—C13—C15	110.09 (19)
O1'—C1'—C1	111.11 (16)	O2—C13—C14	111.1 (2)
O1'—C1'—H1'A	109.4	O1—C13—C14	107.84 (18)
C1—C1'—H1'A	109.4	C15—C13—C14	113.2 (2)

O1'—C1'—H1'B	109.4	C13—C14—H14A	109.5
C1—C1'—H1'B	109.4	C13—C14—H14B	109.5
H1'A—C1'—H1'B	108.0	H14A—C14—H14B	109.5
O2—C2—C3	110.05 (14)	C13—C14—H14C	109.5
O2—C2—C1	103.47 (16)	H14A—C14—H14C	109.5
C3—C2—C1	103.91 (14)	H14B—C14—H14C	109.5
O2—C2—H2	112.9	C13—C15—H15A	109.5
C3—C2—H2	112.9	C13—C15—H15B	109.5
C1—C2—H2	112.9	H15A—C15—H15B	109.5
O3—C3—C4	109.35 (14)	C13—C15—H15C	109.5
O3—C3—C2	109.03 (15)	H15A—C15—H15C	109.5
C4—C3—C2	100.96 (15)	H15B—C15—H15C	109.5
O3—C3—H3	112.3	C21—C16—C17	120.91 (18)
C4—C3—H3	112.3	C21—C16—S1	119.27 (15)
C2—C3—H3	112.3	C17—C16—S1	119.78 (16)
O4—C4—C5	108.83 (15)	C18—C17—C16	119.0 (2)
O4—C4—C3	104.34 (13)	C18—C17—H17	120.5
C5—C4—C3	113.49 (16)	C16—C17—H17	120.5
O4—C4—H4	110.0	C17—C18—C19	121.5 (2)
C5—C4—H4	110.0	C17—C18—H18	119.3
C3—C4—H4	110.0	C19—C18—H18	119.3
O5—C5—C4	106.54 (15)	C18—C19—C20	118.72 (19)
O5—C5—H5A	110.4	C18—C19—C22	120.9 (2)
C4—C5—H5A	110.4	C20—C19—C22	120.3 (2)
O5—C5—H5B	110.4	C19—C20—C21	120.8 (2)
C4—C5—H5B	110.4	C19—C20—H20	119.6
H5A—C5—H5B	108.6	C21—C20—H20	119.6
O1'—C6—C7	109.9 (2)	C16—C21—C20	119.0 (2)
O1'—C6—H6A	109.7	C16—C21—H21	120.5
C7—C6—H6A	109.7	C20—C21—H21	120.5
O1'—C6—H6B	109.7	C19—C22—H22A	109.5
C7—C6—H6B	109.7	C19—C22—H22B	109.5
H6A—C6—H6B	108.2	H22A—C22—H22B	109.5
C12—C7—C8	119.1 (2)	C19—C22—H22C	109.5
C12—C7—C6	121.6 (2)	H22A—C22—H22C	109.5
C8—C7—C6	119.3 (2)	H22B—C22—H22C	109.5
O6—S1—O5—C5	178.98 (14)	C1'—O1'—C6—C7	176.95 (18)
O7—S1—O5—C5	48.72 (15)	O1'—C6—C7—C12	-22.5 (3)
C16—S1—O5—C5	-66.49 (15)	O1'—C6—C7—C8	157.6 (2)
C13—O1—C1—O4	103.77 (18)	C12—C7—C8—C9	-0.8 (4)
C13—O1—C1—C1'	-138.45 (17)	C6—C7—C8—C9	179.1 (2)
C13—O1—C1—C2	-11.3 (2)	C7—C8—C9—C10	0.6 (4)
C4—O4—C1—O1	-106.51 (17)	C8—C9—C10—C11	0.0 (4)
C4—O4—C1—C1'	133.06 (17)	C9—C10—C11—C12	-0.4 (4)
C4—O4—C1—C2	7.94 (19)	C8—C7—C12—C11	0.3 (4)
C6—O1'—C1'—C1	-109.0 (2)	C6—C7—C12—C11	-179.6 (2)
O1—C1—C1'—O1'	65.6 (2)	C10—C11—C12—C7	0.3 (4)
O4—C1—C1'—O1'	-173.29 (19)	C2—O2—C13—O1	16.1 (2)

C2—C1—C1'—O1'	−54.9 (3)	C2—O2—C13—C15	134.23 (18)
C13—O2—C2—C3	−132.90 (17)	C2—O2—C13—C14	−100.7 (2)
C13—O2—C2—C1	−22.38 (18)	C1—O1—C13—O2	−2.2 (2)
O1—C1—C2—O2	20.33 (18)	C1—O1—C13—C15	−119.2 (2)
O4—C1—C2—O2	−98.26 (16)	C1—O1—C13—C14	116.9 (2)
C1'—C1—C2—O2	143.52 (16)	O6—S1—C16—C21	−144.04 (16)
O1—C1—C2—C3	135.33 (15)	O7—S1—C16—C21	−12.19 (18)
O4—C1—C2—C3	16.74 (18)	O5—S1—C16—C21	104.34 (15)
C1'—C1—C2—C3	−101.48 (18)	O6—S1—C16—C17	33.81 (17)
O2—C2—C3—O3	−167.94 (15)	O7—S1—C16—C17	165.66 (15)
C1—C2—C3—O3	81.83 (18)	O5—S1—C16—C17	−77.81 (17)
O2—C2—C3—C4	76.98 (18)	C21—C16—C17—C18	2.5 (3)
C1—C2—C3—C4	−33.25 (16)	S1—C16—C17—C18	−175.36 (15)
C1—O4—C4—C5	−151.21 (16)	C16—C17—C18—C19	−0.6 (3)
C1—O4—C4—C3	−29.75 (18)	C17—C18—C19—C20	−2.0 (3)
O3—C3—C4—O4	−76.32 (17)	C17—C18—C19—C22	176.73 (19)
C2—C3—C4—O4	38.51 (16)	C18—C19—C20—C21	2.9 (3)
O3—C3—C4—C5	42.0 (2)	C22—C19—C20—C21	−175.87 (18)
C2—C3—C4—C5	156.83 (15)	C17—C16—C21—C20	−1.6 (3)
S1—O5—C5—C4	165.40 (12)	S1—C16—C21—C20	176.22 (14)
O4—C4—C5—O5	−62.69 (18)	C19—C20—C21—C16	−1.1 (3)
C3—C4—C5—O5	−178.38 (15)		

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H···A
O3—H3O···O4 ⁱ	0.84	1.98	2.812 (2)	174

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