

Ranunculin

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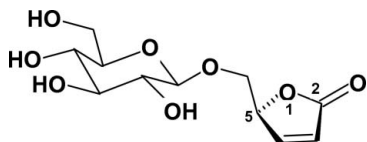
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 Key indicators: single-crystal X-ray study; $T = 173$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; R factor = 0.026; wR factor = 0.069; data-to-parameter ratio = 6.2.

In the title molecule {systematic name: (5*S*)-5-[(β -D-glucopyranosyloxy)methyl]furan-2(*5H*)-one}, $\text{C}_{11}\text{H}_{16}\text{O}_8$, the five-membered ring is essentially planar, the maximum deviation being 0.0151 (13) Å for the O atom. The six-membered ring adopts a chair conformation with puckering parameters $Q = 0.581$ (2) Å, $\theta = 9.0$ (2)° and $\varphi = 39.7$ (13)°, and with all of the substituents of the glucoside unit having normal equatorial orientations. The crystal structure is stabilized by extensive O—H...O and C—H...O hydrogen bonding, resulting in a three-dimensional network.

Related literature

For background to ranunculin, see: Hill & van Heyningen (1951); Bai *et al.* (1996); Benn & Yelland (1968); Boll (1968); Camps *et al.* (1982); Fang *et al.* (1989). For chemical and spectrometric data for closely related, simple butenolides, see: Perry *et al.* (1996). For comparison bond distances, see: Allen *et al.* (1987). For puckering parameters, see: Cremer & Pople (1975).



Experimental

Crystal data

$\text{C}_{11}\text{H}_{16}\text{O}_8$
 $M_r = 276.24$
 Monoclinic, $P2_1$
 $a = 5.7944$ (4) Å
 $b = 6.9359$ (3) Å
 $c = 15.0491$ (10) Å
 $\beta = 97.895$ (2)°

$V = 599.08$ (6) Å³
 $Z = 2$
 Mo $K\alpha$ radiation
 $\mu = 0.13$ mm⁻¹
 $T = 173$ K
 $0.30 \times 0.24 \times 0.02$ mm

Data collection

Nonius KappaCCD diffractometer with APEXII CCD
 Absorption correction: multi-scan (SORTAV; Blessing, 1997)
 $T_{\min} = 0.961$, $T_{\max} = 0.997$
 1926 measured reflections
 1133 independent reflections
 1112 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.018$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.026$
 $wR(F^2) = 0.069$
 $S = 1.04$
 1133 reflections
 184 parameters
 1 restraint

H atoms treated by a mixture of independent and constrained refinement

$\Delta\rho_{\max} = 0.18$ e Å⁻³
 $\Delta\rho_{\min} = -0.16$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

D—H...A	D—H	H...A	D...A	D—H...A
O5—H5O...O8 ⁱ	0.86 (3)	2.17 (3)	2.910 (2)	144 (3)
O6—H6O...O5 ⁱⁱ	0.92 (3)	1.94 (3)	2.824 (2)	162 (3)
O7—H7O...O6 ⁱⁱⁱ	0.84 (3)	1.84 (3)	2.668 (2)	173 (3)
O8—H8O...O2 ⁱⁱⁱ	0.88 (3)	1.96 (3)	2.830 (2)	167 (3)
O6—H6O...O7	0.92 (3)	2.56 (3)	2.894 (2)	102 (2)

Symmetry codes: (i) $x - 1, y - 1, z$; (ii) $-x, y + \frac{1}{2}, -z + 1$; (iii) $-x + 2, y + \frac{1}{2}, -z + 2$.

Data collection: COLLECT (Hooft, 1998); cell refinement: DENZO (Otwinowski & Minor, 1997); data reduction: SCALEPACK (Otwinowski & Minor, 1997); program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: ORTEP-3 for Windows (Farrugia, 1997); software used to prepare material for publication: SHELXL97.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: FL2315).

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supporting information

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Michael Benn, Lois Jean Yelland and Masood Parvez

S1. Comment

Ranunculin (I) is the glycosidic precursor of the vesicant protoanemonin present in numerous species of *Ranunculaceae* and is especially associated with the burning sensation on chewing leaves of buttercup plants. It was first obtained in crystalline form by Hill and van Heyningen (1951) who established its gross structure and showed that it undergoes enzymatic cleavage by β -glucosidase to yield the aglucone, which underwent easy dehydration to protoanemonin. These processes were shown to occur readily under autolytic conditions (Bai *et al.*, 1996). The *S*-stereochemistry of the dihydrofuranone ring was deduced by Benn and Yelland (1968), and Boll (1968), as shown in the schematic diagram, and later confirmed by synthesis (Camps *et al.*, 1982; Fang *et al.*, 1989). Our sample of (I) was a natural product, and as such had been biosynthesised in the plant (and not made in a laboratory). The only stereoisomer which occurs naturally is the *D*-isomer, and both that and the anomeric configuration of the glycosidic bond in (I) were established by cleavage of the glycoside by β -D-glucosidases. The X-ray structure reported here provides the simplest unequivocal proof of that stereochemistry since the chirality follows from that of the *D*-glucopyranosyl moiety.

The molecular structure of (I) is presented in Fig. 1. The five membered ring, O1/C2—C5, is essentially planar with the maximum deviation of any atom being 0.0151 (13) Å for O1. The six-membered ring adopts a chair conformation with puckering parameters (Cremer & Pople, 1975): $Q = 0.581$ (2) Å, $\theta = 9.0$ (2)° and $\varphi = 39.7$ (13)°, with all of the substituents of the glucoside unit having normal equatorial orientations. The 5-membered ring folds back away from the six-membered ring as reflected by the torsion angle C6—O3—C1—C5 -162.86 (17)°. The bond distances and angles are as expected (Allen, 2002). The structure is stabilized by extensive O—H...O and C—H...O hydrogen bonding resulting in a three dimensional network (Table 1 & Fig. 2).

There are two other, closely related, simple butenolides known, though their structures rest on chemical and spectrometric data, i.e., without X-ray crystallographic support. They are the (*5R,6R*) and (*5S,6R*) stereoisomers of 5-([1- β -D-glucopyranosyloxy]ethyl)-2(*5H*)-furanone (Perry *et al.*, 1996); the glycosidic precursor of (*Z*)-5-ethylidene-2(*5H*)-furanone, a homologue of protoanemonin in *Halocarpus biformis juvenis* foliage.

S2. Experimental

The details of the isolation, and some of the physical properties, of our sample of (I) have been reported previously (Benn & Yelland 1968). Suitable crystals of the title compound for X-ray study were grown from a solution in aqueous ethanol (*ca* 1:20) in the form of plates.

S3. Refinement

An absolute structure could not be established reliably because of insufficient anomalous scattering effects. Therefore, 792 Friedel pairs were merged. The H-atoms were located from difference maps and were included in the refinements at geometrically idealized positions with C—H distances = 0.95, 0.99 and 1.00 Å for aryl, methylene and methine type H-

atoms, respectively; the positions of hydroxyl H-atoms were allowed to refine freely. The H-atoms were assigned $U_{\text{iso}} = 1.2$ and $1.5 \times U_{\text{eq}}$ of the parent C and O-atoms, respectively. The final difference map was free of chemically significant features.

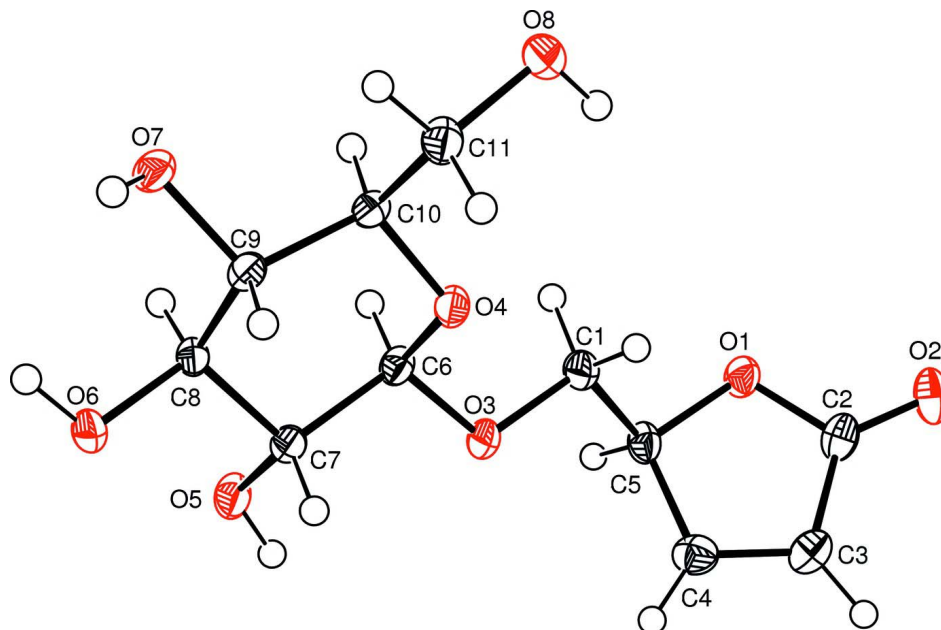


Figure 1

ORTEP-3 (Farrugia, 1997) drawing of the title compound with displacement ellipsoids plotted at 50% probability level.

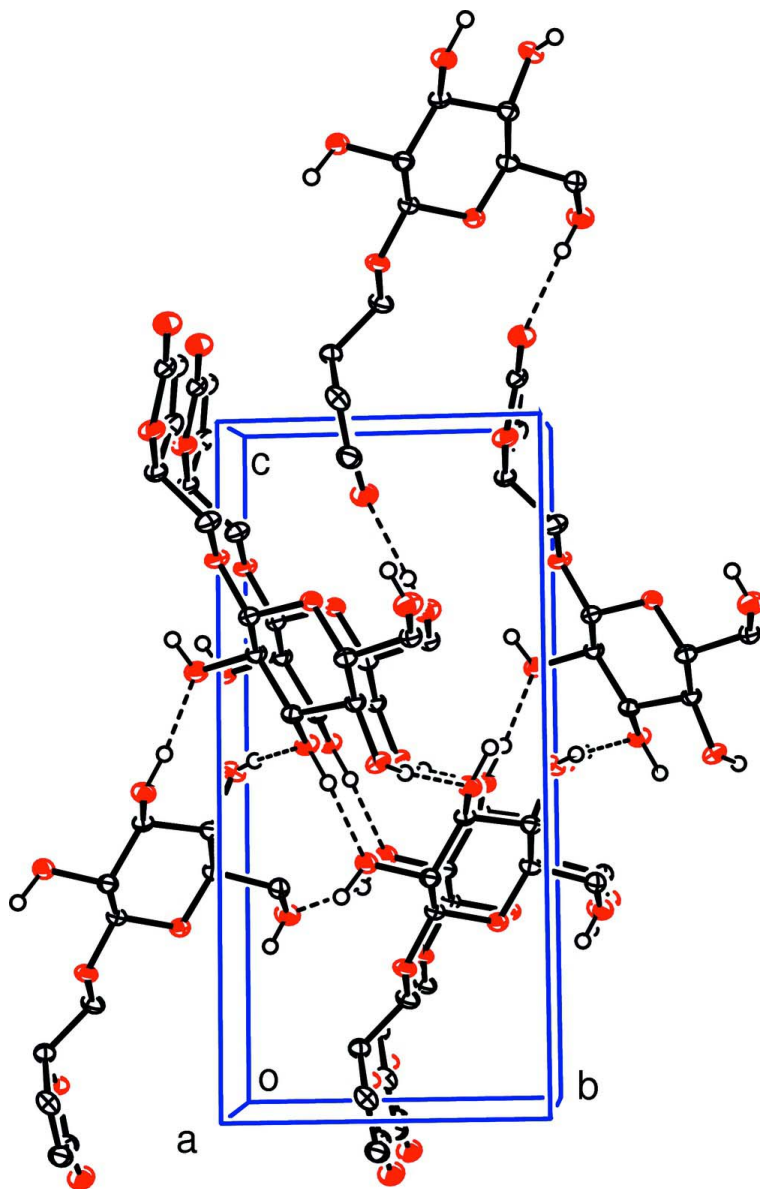


Figure 2

Unit cell packing of the title compound showing intermolecular hydrogen bonds of O—H \cdots O; H-atoms not involved in hydrogen bonds have been excluded.

(5*S*)-5-[(β -D-glucopyranosyloxy)methyl]furan-2(5*H*)-one

Crystal data

C₁₁H₁₆O₈

$M_r = 276.24$

Monoclinic, $P2_1$

Hall symbol: P 2yb

$a = 5.7944 (4) \text{ \AA}$

$b = 6.9359 (3) \text{ \AA}$

$c = 15.0491 (10) \text{ \AA}$

$\beta = 97.895 (2)^\circ$

$V = 599.08 (6) \text{ \AA}^3$

$Z = 2$

$F(000) = 292$

$D_x = 1.531 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$

Cell parameters from 1560 reflections

$\theta = 1.0\text{--}30.0^\circ$

$\mu = 0.13 \text{ mm}^{-1}$

$T = 173$ K
Plate, colorless

$0.30 \times 0.24 \times 0.02$ mm

Data collection

Nonius APEXII CCD [APEXII is a Bruker machine - is this a KappaCCD upgraded with an APEXII CCD?] diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
 φ & ω scans
Absorption correction: multi-scan (SORTAV; Blessing, 1997)

$T_{\min} = 0.961$, $T_{\max} = 0.997$
1926 measured reflections
1133 independent reflections
1112 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.018$
 $\theta_{\max} = 25.0^\circ$, $\theta_{\min} = 3.2^\circ$
 $h = -6 \rightarrow 6$
 $k = -6 \rightarrow 8$
 $l = -17 \rightarrow 17$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.026$
 $wR(F^2) = 0.069$
 $S = 1.04$
1133 reflections
184 parameters
1 restraint
Primary atom site location: structure-invariant direct methods

Secondary atom site location: difference Fourier map
Hydrogen site location: difference Fourier map
H atoms treated by a mixture of independent and constrained refinement
 $w = 1/[\sigma^2(F_o^2) + (0.040P)^2 + 0.1617P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.002$
 $\Delta\rho_{\max} = 0.18 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.16 \text{ e } \text{\AA}^{-3}$

Special details

Experimental. NMR data (400 MHz, ^1H ; 100 MHz ^{13}C) for a solution in D_2O containing sodium 3-trimethylsilylpropionate-2,3 - d_4 as reference: δ_{H} (400 MHz) 7.77 (1H, dd, $J = 1.5$ and 5.8 Hz, H-4), 6.3 (1H, dd, $J = 2.1$ and 5.8 Hz, H-3), 5.47 (1H, m), 4.48 (1H, d, $J = 7.9$ Hz, H-1'), 4.30 (1H, dd, $J = 3.2$ and 12.2 Hz, H-6 A), 3.95 (1H, dd, $J = 5.8$ and 12.2 Hz, H-6B), 3.91 (1H, dd, $J = 2.1$ and 12.5 Hz, H-6A'), 3.72 (1H, dd, $J = 5.8$ and 12.5 Hz, H-6B'), 3.48 (1H, dd, dd, $J = ca$ 9 Hz H-3'), 3.43 (1H, m, H-5'), 3.37 (1H, dd, $J = ca$ 9 Hz, H=4'), and 3.25 (1H, dd, $J = 7.9$ and 9.2 Hz, H-2'); δ_{C} 179.2 s (C-2), 158.5 d (C-4), 124.7 d (C-3), 105.6 d (C-1'), 86.9 d (C-5), 78.7 d (C-5'), 78.3 d (C-3'), 75.6 d (C-2'), 72.2 d (C-4'), 71.7 t (C-6), 63.3 t (C-6').

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 F^2 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)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.9146 (3)	-0.1112 (2)	0.96815 (9)	0.0225 (4)
O2	1.0618 (3)	-0.0493 (3)	1.11025 (10)	0.0309 (4)
O3	0.4499 (2)	0.0377 (2)	0.79524 (9)	0.0200 (3)
O4	0.5209 (2)	0.3269 (2)	0.73224 (9)	0.0188 (3)
O5	0.1227 (3)	-0.0568 (2)	0.63887 (10)	0.0241 (4)
H5O	0.085 (5)	-0.121 (5)	0.6833 (19)	0.036*
O6	-0.0241 (2)	0.2692 (2)	0.53178 (10)	0.0208 (4)

H6O	-0.025 (5)	0.336 (5)	0.479 (2)	0.031*
O7	0.3667 (3)	0.5126 (3)	0.50561 (9)	0.0215 (3)
H7O	0.259 (5)	0.591 (5)	0.4895 (18)	0.032*
O8	0.8805 (3)	0.6376 (3)	0.72096 (10)	0.0277 (4)
H8O	0.893 (5)	0.563 (5)	0.769 (2)	0.042*
C1	0.6798 (3)	0.0332 (4)	0.84401 (13)	0.0216 (5)
H1A	0.7191	0.1601	0.8721	0.026*
H1B	0.7947	0.0036	0.8030	0.026*
C2	0.8918 (4)	-0.0724 (3)	1.05518 (13)	0.0225 (5)
C3	0.6434 (4)	-0.0666 (4)	1.06406 (14)	0.0252 (5)
H3	0.5794	-0.0486	1.1183	0.030*
C4	0.5229 (4)	-0.0912 (3)	0.98301 (14)	0.0243 (5)
H4	0.3578	-0.0895	0.9699	0.029*
C5	0.6862 (4)	-0.1216 (3)	0.91570 (13)	0.0204 (5)
H5	0.6609	-0.2517	0.8874	0.025*
C6	0.4469 (4)	0.1343 (3)	0.71336 (13)	0.0178 (4)
H6	0.5552	0.0694	0.6765	0.021*
C7	0.2004 (3)	0.1338 (3)	0.66293 (13)	0.0176 (4)
H7	0.0906	0.1980	0.6994	0.021*
C8	0.2093 (3)	0.2433 (3)	0.57590 (13)	0.0167 (4)
H8	0.2985	0.1652	0.5363	0.020*
C9	0.3238 (3)	0.4413 (3)	0.59065 (13)	0.0172 (4)
H9	0.2164	0.5310	0.6170	0.021*
C10	0.5559 (3)	0.4305 (3)	0.65204 (13)	0.0180 (4)
H10	0.6738	0.3622	0.6208	0.022*
C11	0.6423 (4)	0.6302 (3)	0.67948 (14)	0.0218 (5)
H11A	0.6248	0.7137	0.6256	0.026*
H11B	0.5424	0.6840	0.7217	0.026*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0203 (7)	0.0294 (9)	0.0169 (7)	0.0038 (6)	-0.0002 (5)	0.0024 (6)
O2	0.0327 (8)	0.0329 (10)	0.0233 (8)	0.0036 (8)	-0.0095 (7)	-0.0014 (8)
O3	0.0212 (7)	0.0242 (8)	0.0137 (7)	-0.0011 (7)	-0.0005 (5)	0.0038 (6)
O4	0.0226 (7)	0.0190 (8)	0.0143 (7)	-0.0026 (6)	0.0005 (5)	-0.0001 (6)
O5	0.0317 (8)	0.0223 (8)	0.0172 (7)	-0.0077 (7)	-0.0007 (6)	0.0020 (7)
O6	0.0177 (7)	0.0259 (8)	0.0174 (7)	-0.0007 (6)	-0.0027 (6)	0.0030 (7)
O7	0.0218 (7)	0.0257 (8)	0.0172 (7)	0.0016 (7)	0.0036 (6)	0.0063 (7)
O8	0.0262 (8)	0.0330 (9)	0.0227 (8)	-0.0102 (8)	-0.0010 (6)	0.0034 (8)
C1	0.0200 (10)	0.0262 (12)	0.0175 (10)	-0.0020 (10)	-0.0010 (8)	0.0041 (9)
C2	0.0302 (11)	0.0188 (10)	0.0175 (9)	0.0063 (10)	0.0000 (9)	0.0026 (9)
C3	0.0306 (11)	0.0257 (11)	0.0201 (10)	0.0037 (10)	0.0064 (8)	0.0025 (10)
C4	0.0235 (10)	0.0234 (12)	0.0264 (12)	0.0004 (10)	0.0044 (9)	0.0071 (10)
C5	0.0217 (10)	0.0214 (11)	0.0168 (10)	-0.0001 (9)	-0.0023 (8)	0.0016 (9)
C6	0.0216 (10)	0.0179 (10)	0.0138 (9)	-0.0003 (9)	0.0020 (7)	0.0016 (9)
C7	0.0190 (10)	0.0188 (10)	0.0151 (9)	-0.0014 (9)	0.0025 (7)	0.0003 (9)
C8	0.0152 (10)	0.0207 (10)	0.0135 (9)	0.0003 (9)	-0.0001 (8)	-0.0008 (8)

C9	0.0185 (10)	0.0191 (10)	0.0143 (9)	0.0010 (9)	0.0038 (7)	0.0010 (9)
C10	0.0180 (10)	0.0216 (10)	0.0147 (9)	-0.0004 (10)	0.0038 (8)	0.0017 (9)
C11	0.0253 (11)	0.0206 (11)	0.0189 (10)	-0.0016 (9)	0.0004 (8)	0.0006 (10)

Geometric parameters (Å, °)

O1—C2	1.361 (2)	C2—C3	1.464 (3)
O1—C5	1.447 (2)	C3—C4	1.331 (3)
O2—C2	1.208 (3)	C3—H3	0.9500
O3—C6	1.401 (2)	C4—C5	1.493 (3)
O3—C1	1.430 (2)	C4—H4	0.9500
O4—C6	1.419 (3)	C5—H5	1.0000
O4—C10	1.443 (2)	C6—C7	1.523 (3)
O5—C7	1.427 (3)	C6—H6	1.0000
O5—H5O	0.86 (3)	C7—C8	1.521 (3)
O6—C8	1.434 (2)	C7—H7	1.0000
O6—H6O	0.92 (3)	C8—C9	1.528 (3)
O7—C9	1.425 (2)	C8—H8	1.0000
O7—H7O	0.84 (3)	C9—C10	1.525 (2)
O8—C11	1.435 (3)	C9—H9	1.0000
O8—H8O	0.88 (3)	C10—C11	1.511 (3)
C1—C5	1.519 (3)	C10—H10	1.0000
C1—H1A	0.9900	C11—H11A	0.9900
C1—H1B	0.9900	C11—H11B	0.9900
C2—O1—C5	109.40 (15)	O4—C6—H6	109.9
C6—O3—C1	111.09 (15)	C7—C6—H6	109.9
C6—O4—C10	112.00 (15)	O5—C7—C8	106.92 (16)
C7—O5—H5O	113 (2)	O5—C7—C6	111.72 (17)
C8—O6—H6O	110.7 (18)	C8—C7—C6	106.69 (16)
C9—O7—H7O	105.7 (19)	O5—C7—H7	110.5
C11—O8—H8O	107 (2)	C8—C7—H7	110.5
O3—C1—C5	108.08 (17)	C6—C7—H7	110.5
O3—C1—H1A	110.1	O6—C8—C7	108.63 (16)
C5—C1—H1A	110.1	O6—C8—C9	108.56 (17)
O3—C1—H1B	110.1	C7—C8—C9	112.90 (16)
C5—C1—H1B	110.1	O6—C8—H8	108.9
H1A—C1—H1B	108.4	C7—C8—H8	108.9
O2—C2—O1	120.56 (19)	C9—C8—H8	108.9
O2—C2—C3	130.8 (2)	O7—C9—C10	108.24 (15)
O1—C2—C3	108.66 (17)	O7—C9—C8	107.89 (16)
C4—C3—C2	108.1 (2)	C10—C9—C8	111.93 (17)
C4—C3—H3	125.9	O7—C9—H9	109.6
C2—C3—H3	125.9	C10—C9—H9	109.6
C3—C4—C5	109.8 (2)	C8—C9—H9	109.6
C3—C4—H4	125.1	O4—C10—C11	107.91 (16)
C5—C4—H4	125.1	O4—C10—C9	108.49 (15)
O1—C5—C4	103.91 (16)	C11—C10—C9	110.63 (17)

O1—C5—C1	106.48 (16)	O4—C10—H10	109.9
C4—C5—C1	115.19 (19)	C11—C10—H10	109.9
O1—C5—H5	110.3	C9—C10—H10	109.9
C4—C5—H5	110.3	O8—C11—C10	114.47 (18)
C1—C5—H5	110.3	O8—C11—H11A	108.6
O3—C6—O4	107.90 (15)	C10—C11—H11A	108.6
O3—C6—C7	109.54 (16)	O8—C11—H11B	108.6
O4—C6—C7	109.80 (16)	C10—C11—H11B	108.6
O3—C6—H6	109.9	H11A—C11—H11B	107.6
C6—O3—C1—C5	-162.86 (17)	O3—C6—C7—C8	179.11 (17)
C5—O1—C2—O2	-176.8 (2)	O4—C6—C7—C8	60.8 (2)
C5—O1—C2—C3	3.2 (2)	O5—C7—C8—O6	68.0 (2)
O2—C2—C3—C4	176.8 (3)	C6—C7—C8—O6	-172.30 (16)
O1—C2—C3—C4	-3.3 (3)	O5—C7—C8—C9	-171.50 (16)
C2—C3—C4—C5	2.0 (3)	C6—C7—C8—C9	-51.8 (2)
C2—O1—C5—C4	-2.0 (2)	O6—C8—C9—O7	-71.42 (19)
C2—O1—C5—C1	120.06 (19)	C7—C8—C9—O7	168.07 (16)
C3—C4—C5—O1	-0.1 (3)	O6—C8—C9—C10	169.60 (14)
C3—C4—C5—C1	-116.2 (2)	C7—C8—C9—C10	49.1 (2)
O3—C1—C5—O1	-175.41 (16)	C6—O4—C10—C11	-177.97 (16)
O3—C1—C5—C4	-60.8 (2)	C6—O4—C10—C9	62.1 (2)
C1—O3—C6—O4	-61.2 (2)	O7—C9—C10—O4	-169.84 (18)
C1—O3—C6—C7	179.26 (17)	C8—C9—C10—O4	-51.1 (2)
C10—O4—C6—O3	171.64 (15)	O7—C9—C10—C11	72.0 (2)
C10—O4—C6—C7	-69.02 (19)	C8—C9—C10—C11	-169.25 (17)
O3—C6—C7—O5	-64.4 (2)	O4—C10—C11—O8	74.2 (2)
O4—C6—C7—O5	177.31 (17)	C9—C10—C11—O8	-167.27 (16)

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
O5—H5O \cdots O8 ⁱ	0.86 (3)	2.17 (3)	2.910 (2)	144 (3)
O6—H6O \cdots O5 ⁱⁱ	0.92 (3)	1.94 (3)	2.824 (2)	162 (3)
O7—H7O \cdots O6 ⁱⁱ	0.84 (3)	1.84 (3)	2.668 (2)	173 (3)
O8—H8O \cdots O2 ⁱⁱⁱ	0.88 (3)	1.96 (3)	2.830 (2)	167 (3)
C3—H3 \cdots O4 ^{iv}	0.95	2.55	3.413 (3)	151
C4—H4 \cdots O1 ^v	0.95	2.57	3.504 (3)	168
C8—H8 \cdots O7 ^{vi}	1.00	2.37	3.306 (3)	155
C1—H1A \cdots O2 ⁱⁱⁱ	0.99	2.38	3.289 (3)	153
C10—H10 \cdots O6 ^{vii}	1.00	2.43	3.415 (3)	167
O6—H6O \cdots O7	0.92 (3)	2.56 (3)	2.894 (2)	102 (2)
C11—H11A \cdots O7	0.99	2.59	2.986 (3)	104

Symmetry codes: (i) $x-1, y-1, z$; (ii) $-x, y+1/2, -z+1$; (iii) $-x+2, y+1/2, -z+2$; (iv) $-x+1, y-1/2, -z+2$; (v) $x-1, y, z$; (vi) $-x+1, y-1/2, -z+1$; (vii) $x+1, y, z$.