

3,3'-Dihydroxy-6,6'-bis(hydroxymethyl)-2,2'-(pentane-1,1-diyl)di-4*H*-pyran-4-one

 Mu-Song Liu,^a Pan-Pan Hu^b and Tao Zhou^{b*}

^aCollege of Pharmaceutical Science, Zhejiang University of Technology, Hangzhou, Zhejiang 310014, People's Republic of China, and ^bSchool of Food Science and Biotechnology, Zhejiang Gongshang University, Hangzhou, Zhejiang 310012, People's Republic of China

Correspondence e-mail: taozhou@zjgsu.edu.cn

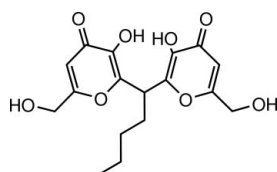
Received 28 February 2012; accepted 8 March 2012

Key indicators: single-crystal X-ray study; $T = 296$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; disorder in main residue; R factor = 0.043; wR factor = 0.120; data-to-parameter ratio = 15.3.

In the title molecule, $\text{C}_{17}\text{H}_{20}\text{O}_8$, the two pyran rings form a dihedral angle of $61.2(2)^\circ$. The two hydroxymethyl groups are each disordered over two sets of sites in a $0.764(3):0.236(3)$ ratio. In the crystal, $\text{O}-\text{H}\cdots\text{O}$ hydrogen bonds link the molecules into layers parallel to the ac plane.

Related literature

For the biological properties of kojic acid, see: Kobayashi *et al.* (1995). For related structures, see: Nurchi *et al.* (2010); Kakkar & Singh (2011); Lokaj *et al.* (1991). For the preparation of the title compound, see: Barham & Nathan Reed (1938).



Experimental

Crystal data

$\text{C}_{17}\text{H}_{20}\text{O}_8$
 $M_r = 352.33$
 Triclinic, $P\bar{1}$
 $a = 6.4234(3)$ Å
 $b = 9.2394(4)$ Å
 $c = 15.8494(7)$ Å
 $\alpha = 79.993(1)^\circ$
 $\beta = 86.689(2)^\circ$

$\gamma = 66.622(1)^\circ$
 $V = 850.22(7)$ Å³
 $Z = 2$
 Mo $K\alpha$ radiation
 $\mu = 0.11$ mm⁻¹
 $T = 296$ K
 $0.49 \times 0.47 \times 0.33$ mm

Data collection

Rigaku R-AXIS RAPID/ZJUG diffractometer
 Absorption correction: multi-scan (*ABSCOR*; Higashi, 1995)
 $T_{\min} = 0.938$, $T_{\max} = 0.965$
 8442 measured reflections
 3840 independent reflections
 2907 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.017$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.043$
 $wR(F^2) = 0.120$
 $S = 1.00$
 3840 reflections
 251 parameters
 6 restraints
 H-atom parameters constrained
 $\Delta\rho_{\max} = 0.30$ e Å⁻³
 $\Delta\rho_{\min} = -0.30$ e Å⁻³

Table 1
 Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{O3}-\text{H3}\cdots\text{O2}^{\text{i}}$	0.82	1.96	2.7275 (15)	155
$\text{O3}-\text{H3}\cdots\text{O2}$	0.82	2.32	2.7440 (15)	113
$\text{O5}-\text{H5}\cdots\text{O6}^{\text{ii}}$	0.82	2.00	2.7488 (16)	151
$\text{O5}-\text{H5}\cdots\text{O6}$	0.82	2.32	2.7436 (16)	113
$\text{O4A}-\text{H4A}\cdots\text{O2}^{\text{iii}}$	0.82	2.07	2.839 (2)	157
$\text{O8A}-\text{H8A}\cdots\text{O6}^{\text{iii}}$	0.82	2.15	2.894 (2)	152

Symmetry codes: (i) $-x, -y + 1, -z + 1$; (ii) $-x, -y + 1, -z$; (iii) $x + 1, y, z$.

Data collection: *PROCESS-AUTO* (Rigaku, 2006); cell refinement: *PROCESS-AUTO*; data reduction: *CrystalStructure* (Rigaku, 2007); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

The work was supported financially by the National Natural Science Foundation of China (grant No. 20972138) and the Qianjiang Scholars Fund, Zhejiang Province (grant No. 2010R10051).

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: CV5255).

References

- Barham, H. N. & Nathan Reed, G. (1938). *J. Am. Chem. Soc.* **60**, 1541–1545.
 Farrugia, L. J. (1997). *J. Appl. Cryst.* **30**, 565.
 Farrugia, L. J. (1999). *J. Appl. Cryst.* **32**, 837–838.
 Higashi, T. (1995). *ABSCOR*. Rigaku Corporation, Tokyo, Japan.
 Kakkar, R. & Singh, C. (2011). *Int. J. Quantum Chem.* **111**, 4318–4329.
 Kobayashi, Y., Kayahara, H., Tadasa, K., Nakamura, T. & Hiroshi, T. (1995). *Biosci. Biotech. Biochem.* **59**, 1745–1746.
 Lokaj, J., Kožisek, J., Koreň, B., Uher, M. & Vrábel, V. (1991). *Acta Cryst.* **C47**, 193–194.
 Nurchi, V. M., Crisponi, G., Lachowicz, J. I., Murgia, S., Pivetta, T., Remelli, M., Rescigno, A., Niclos-Gutierrez, J., Gonzalez-Perez, J. M., Dominguez-Martin, A., Castineiras, A. & Szewczuk, K. (2010). *J. Inorg. Biochem.* **104**, 560–569.
 Rigaku (2006). *PROCESS-AUTO*. Rigaku Corporation, Tokyo, Japan.
 Rigaku (2007). *CrystalStructure*. Rigaku Corporation, Tokyo, Japan.
 Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112–122.

supplementary materials

Acta Cryst. (2012). E68, o1064 [doi:10.1107/S1600536812010276]

3,3'-Dihydroxy-6,6'-bis(hydroxymethyl)-2,2'-(pentane-1,1-diyl)di-4*H*-pyran-4-one

Mu-Song Liu, Pan-Pan Hu and Tao Zhou

Comment

Kojic acid possesses appreciable inhibitory activity against tyrosinase, a key enzyme in the biosynthesis of melanin, due to its ability to chelate copper in the active site of this enzyme. Thus kojic acid inhibits the production of melanin pigment, consequently being used in cosmetic. Therefore, in an attempt to seek potent tyrosinase inhibitors, the derivatives of kojic acid have been widely investigated (Kobayashi *et al.*, 1995). There were little attention to crystal structure of kojic and derivatives. The similar crystal structure of Kojic acid have been reported on 5-hydroxy-2-(hydroxymethyl)-4*H*-pyran-one (Lokaj *et al.*, 1991; Kakkar *et al.*, 2011) and 6,6'-methylenebis (5-hydroxy-2-(hydroxymethyl)-4*H*-pyran-4-one) (Nurchi *et al.*, 2010). Herein, we report the crystal structure of the title compound (I).

In (I) (Fig. 1), two pyranone rings are planar forming the dihedral angle of 61.2 (2)°. Hydroxyl groups are almost coplanar with their linked pyranone rings forming the torsion angles O2—C3—C4—O3 of -0.10° and O6—C10—C9—O5 of 1.24°.

Intermolecular hydrogen bonds O3—H3⋯O2ⁱ and O5—H5⋯O6ⁱⁱ (Table 1) link molecules into zigzag chains along the *c* axis. Further, intermolecular hydrogen bonds O4A—H4⋯O2ⁱⁱⁱ and O8A—H8A⋯O6ⁱⁱⁱ (Table 1) link all of the components of the structure into layers parallel to *ac* plane.

Experimental

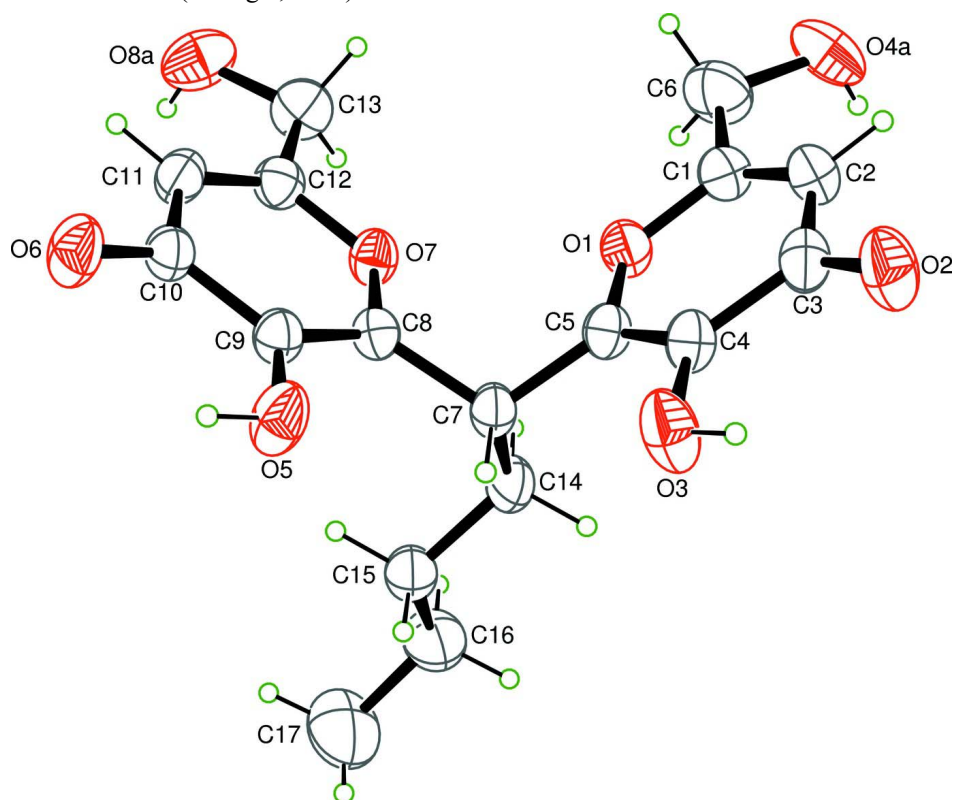
To a solution of 5-hydroxy-(2-hydroxymethyl)-4*H*-pyran-4-one (kojic acid) (1.42 g, 10 mmol), sodium carbonate (1.06 g, 10 mmol) in water (10 ml) and methanol (10 ml) was added pentanal (10 mmol) at 343k with stirring. The stirring was continued for 3 h at that temperature. After removal of about half volume of the solvent, the solution was neutralized to pH=1 with concentrated hydrochloride. The crude product was obtained by filtration as an off-white solid (1.32, 75%), which was recrystallized from dichloromethane solution, giving colorless crystals of the title compound suitable for X-ray diffraction. ¹H NMR (400 MHz DMSO) / d 0.84 (t, J = 6.8 Hz, 3H), 1.21 (m, 4H), 1.29 (2, m), 1.92 (m, 2H), 4.28 (s, 4H), 4.68 (t, J = 8.0 Hz, 1H), 5.62 (s, 2H), 6.28 (s, 2H), 9.03 (s, 2H).

Refinement

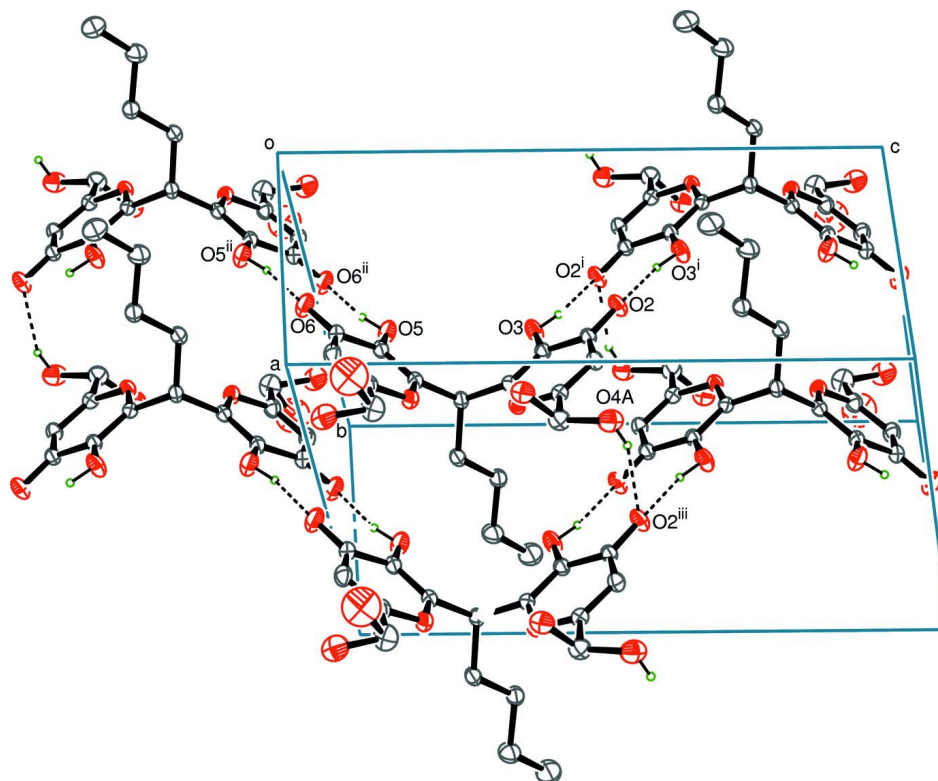
H atoms were placed in calculated positions with O—H = 0.82 and C—H = 0.93–0.98 Å, and included in the refinement in riding model, with $U_{\text{iso}}(\text{H}) = 1.2 - 1.5 U_{\text{eq}}(\text{carrier atom})$. Atoms O4 and O8 of hydroxyl groups were treated as disordered over two positions - A and B, respectively - with the occupancies refined to 0.764 (3) and 0.236 (3), respectively.

Computing details

Data collection: *PROCESS-AUTO* (Rigaku, 2006); cell refinement: *PROCESS-AUTO* (Rigaku, 2006); data reduction: *CrystalStructure* (Rigaku, 2007); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

**Figure 1**

Molecular structure of (I) showing atom-labelling scheme and displacement ellipsoids at 50% probability level. Only major components of the disordered groups are shown.


Figure 2

A portion of the crystal packing of (I) with hydrogen bonds shown by dashed lines [symmetry codes: (i) $-x, 1 - y, -z$; (ii) $x, 1 - y, 1 - z$; (iii) $1 + x, y, z$]. H atoms not involved in hydrogen bonding have been omitted for clarity.

3,3'-Dihydroxy-6,6'-bis(hydroxymethyl)-2,2'-(pentane-1,1-diyl)di-4*H*-pyran-4-one

Crystal data

$C_{17}H_{20}O_8$
 $M_r = 352.33$
 Triclinic, $P\bar{1}$
 Hall symbol: $-P\ 1$
 $a = 6.4234\ (3)\ \text{\AA}$
 $b = 9.2394\ (4)\ \text{\AA}$
 $c = 15.8494\ (7)\ \text{\AA}$
 $\alpha = 79.993\ (1)^\circ$
 $\beta = 86.689\ (2)^\circ$
 $\gamma = 66.622\ (1)^\circ$
 $V = 850.22\ (7)\ \text{\AA}^3$

$Z = 2$
 $F(000) = 372$
 $D_x = 1.376\ \text{Mg m}^{-3}$
 Mo $K\alpha$ radiation, $\lambda = 0.71073\ \text{\AA}$
 Cell parameters from 6451 reflections
 $\theta = 3.3\text{--}27.4^\circ$
 $\mu = 0.11\ \text{mm}^{-1}$
 $T = 296\ \text{K}$
 Chunk, yellow
 $0.49 \times 0.47 \times 0.33\ \text{mm}$

Data collection

Rigaku R-AXIS RAPID/ZJUG
 diffractometer
 Radiation source: rotating anode
 Graphite monochromator
 Detector resolution: $10.00\ \text{pixels mm}^{-1}$
 ω scans
 Absorption correction: multi-scan
 (ABSCOR; Higashi, 1995)
 $T_{\min} = 0.938$, $T_{\max} = 0.965$

8442 measured reflections
 3840 independent reflections
 2907 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.017$
 $\theta_{\max} = 27.4^\circ$, $\theta_{\min} = 3.3^\circ$
 $h = -8 \rightarrow 8$
 $k = -11 \rightarrow 10$
 $l = -20 \rightarrow 20$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.043$

$wR(F^2) = 0.120$

$S = 1.00$

3840 reflections

251 parameters

6 restraints

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

$w = 1/[\sigma^2(F_o^2) + (0.0554P)^2 + 0.2871P]$

where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} < 0.001$

$\Delta\rho_{\max} = 0.30 \text{ e } \text{\AA}^{-3}$

$\Delta\rho_{\min} = -0.30 \text{ e } \text{\AA}^{-3}$

Special details

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	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
O1	0.70599 (16)	0.37814 (12)	0.33797 (6)	0.0343 (2)	
O2	0.24351 (19)	0.37234 (15)	0.52086 (7)	0.0489 (3)	
O3	0.10419 (18)	0.56659 (15)	0.36549 (7)	0.0499 (3)	
H3	0.0303	0.5617	0.4091	0.075*	
O5	0.10800 (19)	0.56285 (16)	0.11434 (8)	0.0534 (3)	
H5	0.0366	0.5594	0.0740	0.080*	
O6	0.2475 (2)	0.35803 (17)	-0.00299 (8)	0.0562 (3)	
O7	0.70230 (17)	0.35409 (12)	0.16422 (6)	0.0381 (3)	
O4A	1.1115 (3)	0.1209 (2)	0.50117 (12)	0.0591 (5)	0.764 (3)
H4A	1.1380	0.1883	0.5209	0.089*	0.764 (3)
O8A	1.1284 (3)	0.0846 (2)	0.04825 (13)	0.0635 (6)	0.764 (3)
H8A	1.1580	0.1543	0.0169	0.095*	0.764 (3)
O4B	1.0634 (8)	0.0558 (6)	0.3693 (3)	0.0474 (14)	0.236 (3)
H4B	1.0161	-0.0084	0.3968	0.071*	0.236 (3)
O8B	1.061 (2)	-0.0030 (16)	0.0999 (9)	0.153 (5)	0.236 (3)
H8B	1.0323	0.0071	0.0490	0.230*	0.236 (3)
C1	0.7734 (2)	0.28416 (17)	0.41491 (9)	0.0348 (3)	
C2	0.6275 (3)	0.27937 (18)	0.47792 (9)	0.0375 (3)	
H2	0.6811	0.2135	0.5301	0.045*	
C3	0.3903 (2)	0.37414 (17)	0.46608 (9)	0.0347 (3)	
C4	0.3252 (2)	0.47378 (17)	0.38243 (9)	0.0329 (3)	
C5	0.4816 (2)	0.47156 (16)	0.32159 (8)	0.0302 (3)	
C6	1.0238 (3)	0.1844 (2)	0.41614 (11)	0.0474 (4)	
H6A	1.0525	0.0971	0.3848	0.057*	0.764 (3)
H6B	1.1022	0.2494	0.3873	0.057*	0.764 (3)
H6C	1.1034	0.2466	0.3877	0.057*	0.236 (3)

H6D	1.0788	0.1461	0.4755	0.057*	0.236 (3)
C7	0.4372 (2)	0.56545 (17)	0.23201 (8)	0.0331 (3)	
H7	0.2759	0.6362	0.2279	0.040*	
C8	0.4808 (2)	0.45696 (17)	0.16640 (8)	0.0332 (3)	
C9	0.3266 (3)	0.46005 (18)	0.11099 (9)	0.0363 (3)	
C10	0.3925 (3)	0.35414 (19)	0.04794 (9)	0.0400 (3)	
C11	0.6259 (3)	0.24898 (19)	0.04981 (10)	0.0437 (4)	
H11	0.6790	0.1780	0.0109	0.052*	
C12	0.7702 (3)	0.25138 (18)	0.10724 (10)	0.0407 (4)	
C13	1.0161 (3)	0.1441 (2)	0.12089 (14)	0.0593 (5)	
H13A	1.0933	0.2027	0.1416	0.071*	0.764 (3)
H13B	1.0285	0.0544	0.1654	0.071*	0.764 (3)
H13C	1.1133	0.1895	0.0883	0.071*	0.236 (3)
H13D	1.0587	0.1215	0.1812	0.071*	0.236 (3)
C14	0.5730 (3)	0.67220 (18)	0.21157 (9)	0.0396 (3)	
H14A	0.7336	0.6051	0.2178	0.047*	
H14B	0.5444	0.7244	0.1523	0.047*	
C15	0.5152 (3)	0.79877 (19)	0.26837 (10)	0.0421 (4)	
H15A	0.5501	0.7469	0.3275	0.050*	
H15B	0.3537	0.8638	0.2640	0.050*	
C16	0.6444 (4)	0.9060 (2)	0.24434 (13)	0.0565 (5)	
H16A	0.8057	0.8408	0.2504	0.068*	
H16B	0.6136	0.9542	0.1845	0.068*	
C17	0.5856 (5)	1.0365 (3)	0.29752 (16)	0.0846 (8)	
H17A	0.4309	1.1097	0.2863	0.127*	
H17B	0.6839	1.0927	0.2830	0.127*	
H17C	0.6046	0.9906	0.3572	0.127*	

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0301 (5)	0.0425 (6)	0.0298 (5)	-0.0143 (4)	0.0034 (4)	-0.0054 (4)
O2	0.0423 (6)	0.0612 (7)	0.0361 (6)	-0.0176 (6)	0.0120 (5)	-0.0004 (5)
O3	0.0323 (6)	0.0651 (8)	0.0361 (6)	-0.0075 (5)	0.0071 (4)	0.0030 (5)
O5	0.0355 (6)	0.0775 (9)	0.0462 (7)	-0.0136 (6)	-0.0004 (5)	-0.0285 (6)
O6	0.0540 (7)	0.0746 (9)	0.0452 (7)	-0.0240 (6)	-0.0054 (5)	-0.0247 (6)
O7	0.0378 (6)	0.0412 (6)	0.0340 (5)	-0.0127 (5)	0.0010 (4)	-0.0101 (4)
O4A	0.0509 (10)	0.0545 (10)	0.0667 (11)	-0.0207 (8)	-0.0180 (8)	0.0091 (8)
O8A	0.0518 (10)	0.0561 (11)	0.0838 (14)	-0.0127 (9)	0.0167 (9)	-0.0404 (10)
O4B	0.038 (3)	0.037 (3)	0.062 (3)	-0.008 (2)	0.009 (2)	-0.014 (2)
O8B	0.153 (5)	0.153 (5)	0.154 (5)	-0.060 (2)	0.0009 (10)	-0.0263 (13)
C1	0.0354 (7)	0.0357 (7)	0.0338 (7)	-0.0140 (6)	-0.0016 (6)	-0.0066 (6)
C2	0.0399 (8)	0.0384 (8)	0.0306 (7)	-0.0136 (6)	-0.0006 (6)	-0.0007 (6)
C3	0.0393 (8)	0.0364 (7)	0.0302 (7)	-0.0170 (6)	0.0054 (6)	-0.0066 (6)
C4	0.0318 (7)	0.0363 (7)	0.0301 (7)	-0.0130 (6)	0.0024 (5)	-0.0058 (6)
C5	0.0312 (7)	0.0334 (7)	0.0282 (6)	-0.0141 (6)	0.0012 (5)	-0.0071 (5)
C6	0.0351 (8)	0.0514 (9)	0.0512 (9)	-0.0134 (7)	-0.0007 (7)	-0.0052 (8)
C7	0.0350 (7)	0.0381 (7)	0.0256 (6)	-0.0136 (6)	0.0025 (5)	-0.0064 (5)
C8	0.0349 (7)	0.0377 (7)	0.0274 (6)	-0.0157 (6)	0.0044 (5)	-0.0046 (6)
C9	0.0369 (8)	0.0450 (8)	0.0283 (7)	-0.0179 (7)	0.0036 (6)	-0.0067 (6)

C10	0.0471 (9)	0.0484 (9)	0.0293 (7)	-0.0235 (7)	0.0023 (6)	-0.0082 (6)
C11	0.0541 (10)	0.0420 (8)	0.0358 (8)	-0.0171 (7)	0.0047 (7)	-0.0140 (7)
C12	0.0460 (9)	0.0372 (8)	0.0362 (8)	-0.0134 (7)	0.0047 (6)	-0.0077 (6)
C13	0.0489 (10)	0.0527 (10)	0.0658 (12)	-0.0064 (8)	0.0031 (9)	-0.0169 (9)
C14	0.0500 (9)	0.0420 (8)	0.0309 (7)	-0.0233 (7)	0.0081 (6)	-0.0063 (6)
C15	0.0520 (9)	0.0409 (8)	0.0358 (8)	-0.0207 (7)	0.0033 (7)	-0.0075 (6)
C16	0.0716 (12)	0.0488 (10)	0.0576 (11)	-0.0329 (9)	0.0096 (9)	-0.0106 (8)
C17	0.129 (2)	0.0774 (15)	0.0786 (15)	-0.0695 (16)	0.0189 (15)	-0.0289 (12)

Geometric parameters (Å, °)

O1—C1	1.3502 (17)	C6—H6B	0.9700
O1—C5	1.3669 (16)	C6—H6C	0.9553
O2—C3	1.2474 (17)	C6—H6D	0.9795
O3—C4	1.3492 (17)	C7—C8	1.5069 (19)
O3—H3	0.8200	C7—C14	1.542 (2)
O5—C9	1.3518 (18)	C7—H7	0.9800
O5—H5	0.8200	C8—C9	1.350 (2)
O6—C10	1.2531 (19)	C9—C10	1.448 (2)
O7—C12	1.3529 (18)	C10—C11	1.426 (2)
O7—C8	1.3641 (17)	C11—C12	1.345 (2)
O4A—C6	1.424 (2)	C11—H11	0.9300
O4A—H4A	0.8200	C12—C13	1.499 (2)
O4A—H6D	0.4491	C13—H13A	0.9700
O8A—C13	1.399 (3)	C13—H13B	0.9700
O8A—H8A	0.8200	C13—H13C	0.9635
O8A—H13C	1.2178	C13—H13D	0.9733
O4B—C6	1.439 (5)	C14—C15	1.516 (2)
O4B—H4B	0.8200	C14—H14A	0.9700
O8B—C13	1.369 (13)	C14—H14B	0.9700
O8B—H8B	0.8200	C15—C16	1.517 (2)
C1—C2	1.337 (2)	C15—H15A	0.9700
C1—C6	1.503 (2)	C15—H15B	0.9700
C2—C3	1.428 (2)	C16—C17	1.504 (3)
C2—H2	0.9300	C16—H16A	0.9700
C3—C4	1.4541 (19)	C16—H16B	0.9700
C4—C5	1.3484 (19)	C17—H17A	0.9600
C5—C7	1.5073 (19)	C17—H17B	0.9600
C6—H6A	0.9700	C17—H17C	0.9600
C1—O1—C5	120.29 (11)	O6—C10—C11	124.42 (15)
C4—O3—H3	109.5	O6—C10—C9	120.11 (15)
C9—O5—H5	109.5	C11—C10—C9	115.47 (13)
C12—O7—C8	120.36 (12)	C12—C11—C10	120.66 (14)
C6—O4A—H4A	109.5	C12—C11—H11	119.7
H4A—O4A—H6D	104.7	C10—C11—H11	119.7
C13—O8A—H8A	109.5	C11—C12—O7	121.98 (14)
H8A—O8A—H13C	70.3	C11—C12—C13	127.48 (15)
C6—O4B—H4B	109.5	O7—C12—C13	110.50 (14)
C13—O8B—H8B	109.5	O8B—C13—O8A	52.8 (6)

C2—C1—O1	122.32 (13)	O8B—C13—C12	111.3 (6)
C2—C1—C6	126.71 (14)	O8A—C13—C12	115.07 (17)
O1—C1—C6	110.89 (13)	O8B—C13—H13A	140.2
C1—C2—C3	120.82 (13)	O8A—C13—H13A	108.5
C1—C2—H2	119.6	C12—C13—H13A	108.5
C3—C2—H2	119.6	O8B—C13—H13B	59.8
O2—C3—C2	124.45 (13)	O8A—C13—H13B	108.5
O2—C3—C4	120.46 (13)	C12—C13—H13B	108.5
C2—C3—C4	115.07 (12)	H13A—C13—H13B	107.5
O3—C4—C5	120.03 (12)	O8B—C13—H13C	108.9
O3—C4—C3	119.06 (12)	O8A—C13—H13C	58.6
C5—C4—C3	120.89 (13)	C12—C13—H13C	112.0
C4—C5—O1	120.60 (12)	H13A—C13—H13C	53.5
C4—C5—C7	126.41 (13)	H13B—C13—H13C	139.0
O1—C5—C7	112.99 (11)	O8B—C13—H13D	104.3
O4A—C6—O4B	109.7 (3)	O8A—C13—H13D	133.4
O4A—C6—C1	111.96 (15)	C12—C13—H13D	111.1
O4B—C6—C1	108.5 (2)	H13A—C13—H13D	60.5
O4A—C6—H6A	109.2	H13B—C13—H13D	48.6
C1—C6—H6A	109.2	H13C—C13—H13D	108.8
O4A—C6—H6B	109.2	C15—C14—C7	113.65 (12)
O4B—C6—H6B	108.1	C15—C14—H14A	108.8
C1—C6—H6B	109.2	C7—C14—H14A	108.8
H6A—C6—H6B	107.9	C15—C14—H14B	108.8
O4A—C6—H6C	109.0	C7—C14—H14B	108.8
O4B—C6—H6C	107.4	H14A—C14—H14B	107.7
C1—C6—H6C	110.2	C14—C15—C16	112.37 (14)
H6A—C6—H6C	107.1	C14—C15—H15A	109.1
O4B—C6—H6D	112.6	C16—C15—H15A	109.1
C1—C6—H6D	109.6	C14—C15—H15B	109.1
H6A—C6—H6D	112.1	C16—C15—H15B	109.1
H6B—C6—H6D	108.8	H15A—C15—H15B	107.9
H6C—C6—H6D	108.6	C17—C16—C15	113.87 (17)
C5—C7—C8	111.20 (12)	C17—C16—H16A	108.8
C5—C7—C14	112.71 (12)	C15—C16—H16A	108.8
C8—C7—C14	110.77 (11)	C17—C16—H16B	108.8
C5—C7—H7	107.3	C15—C16—H16B	108.8
C8—C7—H7	107.3	H16A—C16—H16B	107.7
C14—C7—H7	107.3	C16—C17—H17A	109.5
C9—C8—O7	120.75 (13)	C16—C17—H17B	109.5
C9—C8—C7	126.05 (13)	H17A—C17—H17B	109.5
O7—C8—C7	113.13 (12)	C16—C17—H17C	109.5
C8—C9—O5	119.76 (13)	H17A—C17—H17C	109.5
C8—C9—C10	120.76 (14)	H17B—C17—H17C	109.5
O5—C9—C10	119.48 (13)		
C5—O1—C1—C2	-0.8 (2)	C5—C7—C8—C9	118.54 (16)
C5—O1—C1—C6	176.06 (12)	C14—C7—C8—C9	-115.30 (16)
O1—C1—C2—C3	0.4 (2)	C5—C7—C8—O7	-64.36 (15)

C6—C1—C2—C3	-175.94 (14)	C14—C7—C8—O7	61.80 (15)
C1—C2—C3—O2	177.73 (16)	O7—C8—C9—O5	179.93 (13)
C1—C2—C3—C4	-0.6 (2)	C7—C8—C9—O5	-3.2 (2)
O2—C3—C4—O3	1.3 (2)	O7—C8—C9—C10	0.1 (2)
C2—C3—C4—O3	179.63 (13)	C7—C8—C9—C10	176.96 (13)
O2—C3—C4—C5	-177.25 (14)	C8—C9—C10—O6	179.85 (15)
C2—C3—C4—C5	1.1 (2)	O5—C9—C10—O6	0.0 (2)
O3—C4—C5—O1	179.94 (13)	C8—C9—C10—C11	0.5 (2)
C3—C4—C5—O1	-1.5 (2)	O5—C9—C10—C11	-179.41 (14)
O3—C4—C5—C7	0.1 (2)	O6—C10—C11—C12	-179.32 (16)
C3—C4—C5—C7	178.57 (13)	C9—C10—C11—C12	0.0 (2)
C1—O1—C5—C4	1.4 (2)	C10—C11—C12—O7	-1.1 (2)
C1—O1—C5—C7	-178.71 (12)	C10—C11—C12—C13	176.58 (17)
C2—C1—C6—O4A	-20.6 (2)	C8—O7—C12—C11	1.7 (2)
O1—C1—C6—O4A	162.71 (14)	C8—O7—C12—C13	-176.37 (14)
C2—C1—C6—O4B	100.7 (3)	C11—C12—C13—O8B	-29.2 (7)
O1—C1—C6—O4B	-76.1 (3)	O7—C12—C13—O8B	148.7 (7)
C4—C5—C7—C8	-111.66 (16)	C11—C12—C13—O8A	28.5 (3)
O1—C5—C7—C8	68.45 (15)	O7—C12—C13—O8A	-153.56 (17)
C4—C5—C7—C14	123.26 (16)	C5—C7—C14—C15	-61.92 (17)
O1—C5—C7—C14	-56.62 (16)	C8—C7—C14—C15	172.77 (13)
C12—O7—C8—C9	-1.1 (2)	C7—C14—C15—C16	-177.67 (14)
C12—O7—C8—C7	-178.40 (12)	C14—C15—C16—C17	178.09 (18)

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

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
O3—H3...O2 ⁱ	0.82	1.96	2.7275 (15)	155
O3—H3...O2	0.82	2.32	2.7440 (15)	113
O5—H5...O6 ⁱⁱ	0.82	2.00	2.7488 (16)	151
O5—H5...O6	0.82	2.32	2.7436 (16)	113
O4A—H4A...O2 ⁱⁱⁱ	0.82	2.07	2.839 (2)	157
O8A—H8A...O6 ⁱⁱⁱ	0.82	2.15	2.894 (2)	152

Symmetry codes: (i) $-x, -y+1, -z+1$; (ii) $-x, -y+1, -z$; (iii) $x+1, y, z$.