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6-Hydroxymethyl-4-methoxy-2H-pyran-2-one (Opuntiol)

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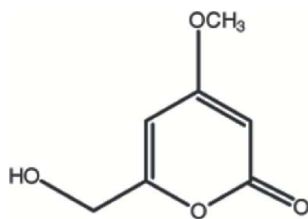
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Key indicators: single-crystal X-ray study; $T = 296$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; R factor = 0.037; wR factor = 0.094; data-to-parameter ratio = 12.1.

The title compound, $\text{C}_7\text{H}_8\text{O}_4$, isolated from *Opuntia dillenii* Haw (Cactaceae), is almost planar [maximum deviation of 0.027 (2) Å] except for the H atoms of the methylene and methyl groups. The crystal packing is stabilized by $\text{C}-\text{H}\cdots\text{O}$ and $\text{O}-\text{H}\cdots\text{O}$ intermolecular hydrogen bonds, resulting in the formation of a three-dimensional network.

Related literature

For the use of the stem and fruit of *Opuntia dillenii* Haw (Cactaceae) in folk medicine, see: Chang *et al.* (2008). For phytochemical investigations of this plant, see: Qiu *et al.* (2002). For comparative bond lengths, see: Allen *et al.* (1987).



Experimental

Crystal data

$\text{C}_7\text{H}_8\text{O}_4$
 $M_r = 156.13$
Monoclinic, $P2_1/c$
 $a = 4.0499$ (5) Å
 $b = 18.101$ (2) Å
 $c = 9.4743$ (13) Å
 $\beta = 96.720$ (7)°

$V = 689.76$ (15) Å³
 $Z = 4$
Mo $K\alpha$ radiation
 $\mu = 0.13$ mm⁻¹
 $T = 296$ K
 $0.34 \times 0.25 \times 0.19$ mm

Data collection

Bruker Kappa APEXII CCD area-detector diffractometer
1268 independent reflections
882 reflections with $I > 2\sigma(I)$
6516 measured reflections
 $R_{\text{int}} = 0.046$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.037$
 $wR(F^2) = 0.094$
 $S = 1.02$
1268 reflections
105 parameters
1 restraint
H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\text{max}} = 0.16$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.16$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{O4}-\text{H1}\cdots\text{O1}^{\text{i}}$	0.836 (14)	2.47 (2)	3.1840 (19)	144.2 (17)
$\text{O4}-\text{H1}\cdots\text{O2}^{\text{i}}$	0.836 (14)	2.073 (13)	2.8678 (18)	158.6 (18)
$\text{C7}-\text{H7B}\cdots\text{O4}^{\text{ii}}$	0.96	2.58	3.494 (2)	159
$\text{C7}-\text{H7C}\cdots\text{O2}^{\text{iii}}$	0.96	2.57	3.504 (2)	164

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

Data collection: APEX2 (Bruker, 2007); cell refinement: SAINT (Bruker, 2007); data reduction: SAINT; program(s) used to solve structure: SIR97 (Altomare *et al.*, 1999); 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) and PLATON (Spek, 2009).

The authors are grateful to the Higher Education Commission for providing financial support. Professor Islam Ullah Khan and Mr Shahzad Sharif are also gratefully acknowledged for providing single-crystal X-ray diffraction facilities at the Materials Chemistry Laboratory, GC University Lahore.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: RK2184).

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6-Hydroxymethyl-4-methoxy-2H-pyran-2-one (Opuntiol)

Muhammad Athar Abbasi, Tayyaba Shahzadi, Mehmet Akkurt, Aziz-ur-Rehman and Tauheeda Riaz

S1. Comment

Opuntia dillenii Haw (Cactaceae) usually grows in semi-desert regions in the tropics and subtropics. The stem and fruit of this plant are used in a folk medicine for reducing cholesterol levels, treatment of gastric ulcers, inflammation, diabetes and several other diseases (Chang *et al.*, 2008). The phytochemical investigations on this plant have led to the isolation of various oxygenated constituents namely *opuntiol*, opuntioside-I, *p*-hydroxybenzoic acid, ethyl 3,4-dihydroxybenzoate, 3,4-dihydroxybenzoic acid, *L*-(-)-malic acid, (*E*)-ferulic acid, 4-ethoxy-6-hydroxymethyl- α -pyrone, 1-heptanecanol, vanillic acid, isorhamnetin, isorhamnetin-3-*O*-rutinoside, rutin, quercetin, 3,3'-dimethyl quercetin, 3-*O*-methyl quercetin, 3-*O*-methyl quercetin 7-*O*- β -*D*-glucopyranoside, kaempferol, kaempferol 7-*O*- β -*D*-glucopyranoside, kaempferol 7-*O*- β -*D*-glucopyranosyl-(1 \rightarrow 4)- β -*D*-glucopyranoside, kaempferide, β -sitosterol, and manghaslin (Qiu *et al.*, 2002). In the present study we first time report the crystal structure of opuntiol using single-crystal XRD.

In the title molecule, **I**, shown in Fig. 1, bond lengths and angles display normal values (Allen *et al.*, 1987). Except the H atoms of the methylene and methyl groups, the molecule of **I** is almost planar, with maximum deviations of 0.027 (2) Å for O4, 0.023 (2) Å for C7, 0.020 (1) Å for O2 and -0.019 (2) Å for C2. For the methoxy and hydroxy methyl groups at the C3 and C5 positions in **I**, the C2–C3–O3–C7 and C4–C3–O3–C7 torsion angles are 0.9 (2)° and -178.94 (15)°, the O1–C5–C6–O4 and C4–C5–C6–O4 torsion angles are 177.77 (14)° and -2.9 (3)°, respectively.

In the crystal structure of **I**, there are non-classical C–H \cdots O and classical O–H \cdots O intermolecular hydrogen bonds (Table 1), forming a three-dimensional network (Figs. 2 and 3).

S2. Experimental

Plant Material: *Opuntia dillenii* Haw (whole plant) was collected from the areas of Mar Balochan, Sangla Hill, Distt. Nankana, Pakistan, in April 2008, and identified by Muhammad Ajaib (Taxonomist), Department of Botany, Government College University, Lahore.

Extraction and isolation: The shade-dried ground whole plant (7 kg) of *Opuntia dillenii* Haw was exhaustively extracted with methanol (10L \times 4) at room temperature. The extract was evaporated to yield the residue (1.1 kg), which was dissolved in distilled water (2.0 L) and partitioned with *n*-hexane (2L \times 4), chloroform (2L \times 4), ethyl acetate (2L \times 4) and *n*-butanol (2L \times 4) respectively. The chloroform soluble extract (157 g) was subjected to column chromatography using hexane with gradient of CHCl₃ and followed by methanol up to 100%. Fifteen fractions (Fr. 1-15) were collected. The Fr. 14 was loaded on flash silica gel and eluted with MeOH : CHCl₃ (2 : 98) to get purified crystals of *opuntiol* (84.7 mg).

S3. Refinement

The H atom of the OH group was located in difference Fourier maps and were refined with a O–H distance restrained to 0.83 (1) Å, with displacement parameters fixed at 1.5 times U_{eq} of the parent O atom. The rest H atoms were placed geometrically, with C–H = 0.93–0.97 Å, and treated using a riding model, with $U_{iso}(H) = 1.2$ or $1.5U_{eq}(\text{parent atom})$.

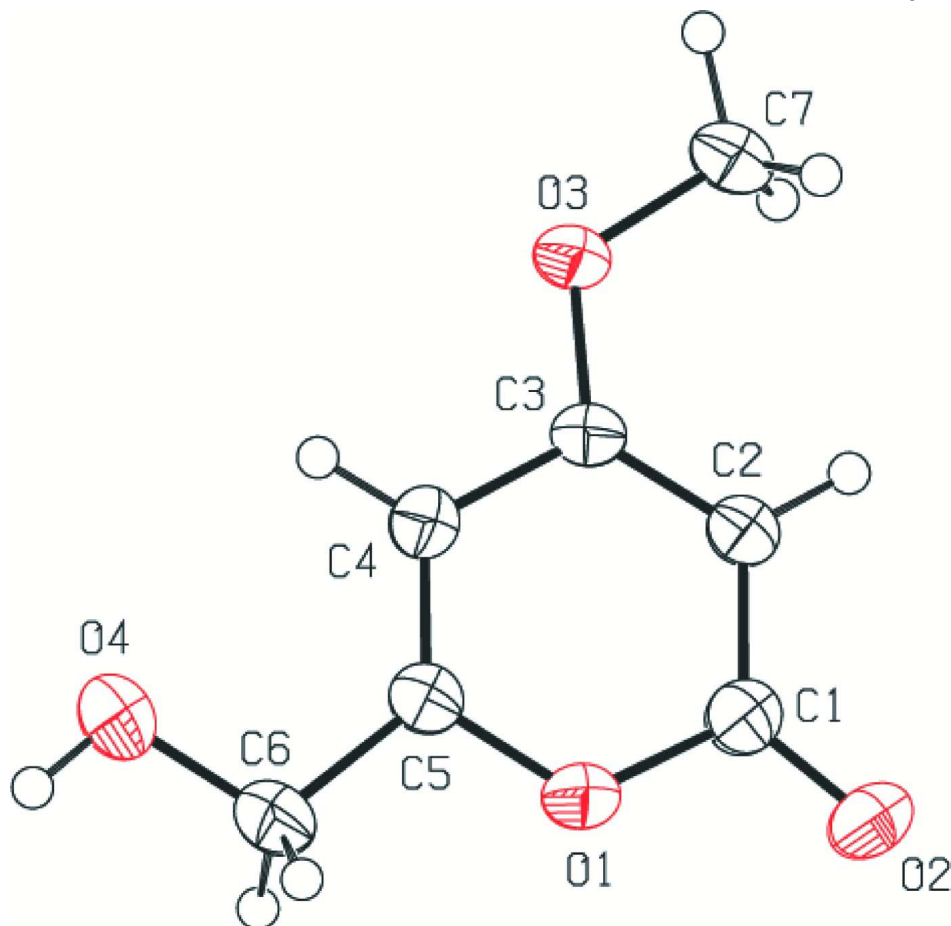
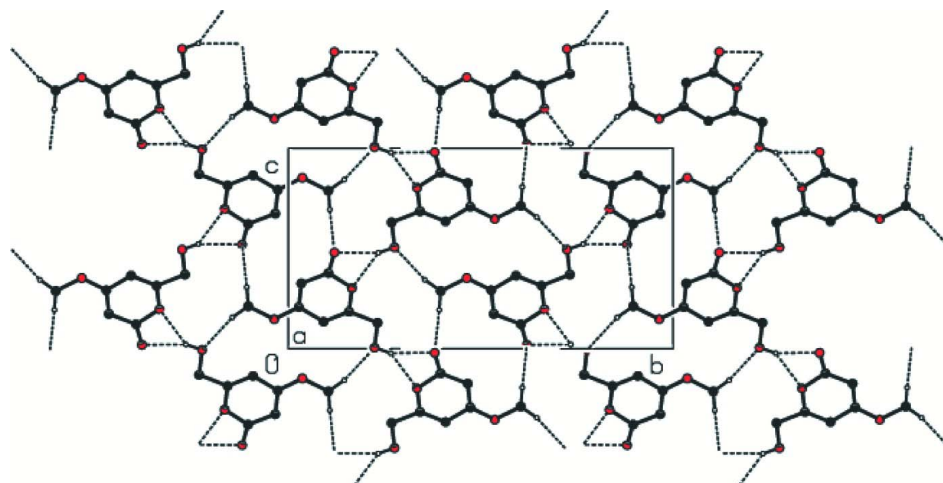
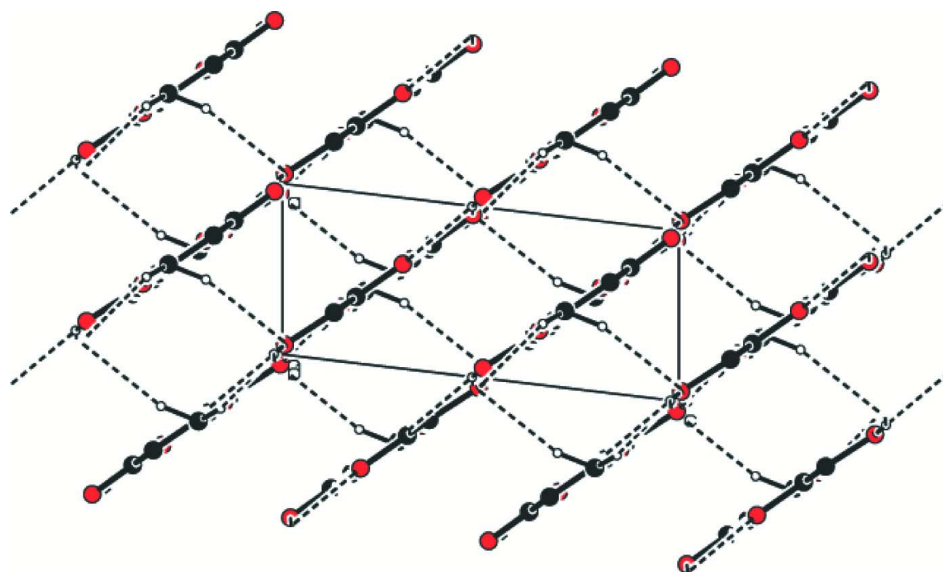


Figure 1

Molecular structure of the title compound with the atom numbering scheme. Displacement ellipsoids for non-H atoms are drawn at the 50% probability level. H atoms are presented as a small spheres of arbitrary radius.

**Figure 2**

The packing and hydrogen bonding of the title compound viewed down *a* axis. Hydrogen atoms not involved in hydrogen bonding have been omitted for clarity.

**Figure 3**

The packing and hydrogen bonding of the title compound viewed down *b* axis. Hydrogen atoms not involved in hydrogen bonding have been omitted for clarity.

6-Hydroxymethyl-4-methoxy-2H-pyran-2-one

Crystal data

$C_7H_8O_4$

$M_r = 156.13$

Monoclinic, $P2_1/c$

Hall symbol: $-P 2_1/c$

$a = 4.0499 (5) \text{ \AA}$

$b = 18.101 (2) \text{ \AA}$

$c = 9.4743 (13) \text{ \AA}$

$\beta = 96.720 (7)^\circ$

$V = 689.76 (15) \text{ \AA}^3$

$Z = 4$

$F(000) = 328$

$D_x = 1.503 \text{ Mg m}^{-3}$

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

Cell parameters from 1306 reflections

$\theta = 2.3\text{--}23.4^\circ$

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

$T = 296$ K $0.34 \times 0.25 \times 0.19$ mm
 Needle, colourless

Data collection

Bruker Kappa APEXII CCD area-detector diffractometer Radiation source: sealed tube Graphite monochromator φ - and ω -scans 6516 measured reflections 1268 independent reflections	882 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.046$ $\theta_{\text{max}} = 25.5^\circ$, $\theta_{\text{min}} = 2.4^\circ$ $h = -4 \rightarrow 4$ $k = -21 \rightarrow 21$ $l = -11 \rightarrow 11$
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Refinement

Refinement on F^2 Least-squares matrix: full $R[F^2 > 2\sigma(F^2)] = 0.037$ $wR(F^2) = 0.094$ $S = 1.02$ 1268 reflections 105 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 atoms treated by a mixture of independent and constrained refinement $w = 1/[\sigma^2(F_o^2) + (0.0406P)^2 + 0.1053P]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\text{max}} < 0.001$ $\Delta\rho_{\text{max}} = 0.16 \text{ e } \text{\AA}^{-3}$ $\Delta\rho_{\text{min}} = -0.16 \text{ e } \text{\AA}^{-3}$ Extinction correction: <i>SHELXL97</i> (Sheldrick, 2008), $FC^* = KFC[1 + 0.001XFC^2\Lambda^3/\text{SIN}(2\Theta)]^{-1/4}$ Extinction coefficient: 0.0080 (19)
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Special details

Geometry. All s.u.'s (except the s.u. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell s.u.'s are taken into account individually in the estimation of s.u.'s in distances, angles and torsion angles; correlations between s.u.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell s.u.'s is used for estimating s.u.'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	0.6125 (3)	0.16542 (6)	0.30332 (12)	0.0376 (4)
O2	0.9490 (3)	0.11978 (7)	0.48000 (14)	0.0495 (5)
O3	0.3187 (3)	-0.03651 (6)	0.14748 (13)	0.0392 (4)
O4	0.0594 (4)	0.22405 (7)	0.00555 (16)	0.0558 (5)
C1	0.7497 (4)	0.10447 (9)	0.3778 (2)	0.0362 (6)
C2	0.6491 (4)	0.03392 (9)	0.32627 (18)	0.0336 (6)
C3	0.4283 (4)	0.02721 (9)	0.20778 (18)	0.0303 (5)
C4	0.2914 (4)	0.09149 (9)	0.13544 (18)	0.0334 (6)
C5	0.3878 (4)	0.15778 (9)	0.18521 (18)	0.0325 (6)
C6	0.2807 (5)	0.23160 (9)	0.1300 (2)	0.0413 (6)
C7	0.4484 (5)	-0.10405 (10)	0.2121 (2)	0.0437 (7)
H1	-0.003 (6)	0.2662 (7)	-0.022 (2)	0.0840*
H2	0.73440	-0.00810	0.37370	0.0400*

H4	0.13750	0.08710	0.05500	0.0400*
H6A	0.17300	0.25800	0.20100	0.0500*
H6B	0.47320	0.25990	0.11000	0.0500*
H7A	0.68640	-0.10410	0.21640	0.0660*
H7B	0.35940	-0.14530	0.15650	0.0660*
H7C	0.38600	-0.10780	0.30650	0.0660*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0434 (7)	0.0266 (7)	0.0405 (8)	-0.0032 (5)	-0.0045 (6)	-0.0040 (6)
O2	0.0590 (9)	0.0408 (8)	0.0437 (8)	-0.0065 (7)	-0.0149 (7)	-0.0057 (7)
O3	0.0503 (8)	0.0224 (7)	0.0420 (8)	-0.0014 (6)	-0.0073 (6)	-0.0020 (6)
O4	0.0688 (10)	0.0331 (8)	0.0596 (10)	0.0072 (7)	-0.0169 (8)	0.0056 (7)
C1	0.0385 (10)	0.0342 (11)	0.0347 (11)	-0.0020 (8)	-0.0002 (8)	0.0000 (9)
C2	0.0385 (10)	0.0256 (10)	0.0355 (11)	-0.0003 (8)	-0.0008 (8)	0.0019 (8)
C3	0.0333 (9)	0.0238 (9)	0.0335 (10)	-0.0023 (7)	0.0026 (8)	-0.0030 (8)
C4	0.0371 (10)	0.0312 (10)	0.0305 (10)	0.0010 (8)	-0.0021 (8)	0.0002 (8)
C5	0.0349 (10)	0.0274 (10)	0.0347 (10)	-0.0006 (8)	0.0022 (8)	0.0007 (8)
C6	0.0459 (11)	0.0267 (10)	0.0500 (12)	0.0014 (8)	0.0000 (9)	0.0015 (9)
C7	0.0548 (12)	0.0240 (10)	0.0501 (13)	0.0006 (8)	-0.0028 (10)	0.0008 (9)

Geometric parameters (Å, °)

O1—C1	1.390 (2)	C4—C5	1.331 (2)
O1—C5	1.364 (2)	C5—C6	1.481 (2)
O2—C1	1.218 (2)	C2—H2	0.9300
O3—C3	1.340 (2)	C4—H4	0.9300
O3—C7	1.439 (2)	C6—H6A	0.9700
O4—C6	1.402 (2)	C6—H6B	0.9700
O4—H1	0.836 (14)	C7—H7A	0.9600
C1—C2	1.410 (2)	C7—H7B	0.9600
C2—C3	1.356 (2)	C7—H7C	0.9600
C3—C4	1.429 (2)		
O1...O4 ⁱ	3.1840 (19)	C2...H7C	2.7800
O2...O4 ⁱ	2.8678 (18)	C2...H7A	2.7200
O2...C6 ⁱ	3.258 (2)	C7...H2	2.5100
O3...C4 ⁱⁱ	3.415 (2)	C7...H7A ^{vii}	3.0900
O4...O2 ⁱⁱⁱ	2.8678 (18)	C7...H6A ^x	2.9900
O4...O1 ⁱⁱⁱ	3.1840 (19)	C7...H6B ^x	2.9800
O1...H1 ⁱ	2.47 (2)	H1...H6B ^{vii}	2.5900
O2...H2 ^{iv}	2.6900	H1...O1 ⁱⁱⁱ	2.47 (2)
O2...H7C ^v	2.5700	H1...O2 ⁱⁱⁱ	2.073 (13)
O2...H1 ⁱ	2.073 (13)	H1...C1 ⁱⁱⁱ	2.676 (15)
O3...H4 ^{vi}	2.6700	H2...C7	2.5100
O4...H7B ^{vi}	2.5800	H2...H7A	2.2800
O4...H4	2.5400	H2...H7C	2.3300

O4...H6B ^{vii}	2.7500	H2...O2 ^{iv}	2.6900
C1...C4 ^{viii}	3.365 (2)	H4...O4	2.5400
C1...C5 ^{viii}	3.470 (2)	H4...O3 ^{vi}	2.6700
C2...C3 ^{viii}	3.473 (2)	H6A...C7 ^{ix}	2.9900
C2...C4 ^{viii}	3.496 (2)	H6B...O4 ^{viii}	2.7500
C3...C2 ^{vii}	3.473 (2)	H6B...H1 ^{viii}	2.5900
C4...C2 ^{vii}	3.496 (2)	H6B...C7 ^{ix}	2.9800
C4...C1 ^{vii}	3.365 (2)	H6B...H7C ^{ix}	2.5700
C4...C7 ⁱⁱ	3.580 (3)	H7A...C2	2.7200
C4...O3 ⁱⁱ	3.415 (2)	H7A...C7 ^{viii}	3.0900
C5...C1 ^{vii}	3.470 (2)	H7A...H2	2.2800
C6...C7 ^{ix}	3.451 (3)	H7B...O4 ^{vi}	2.5800
C6...O2 ⁱⁱⁱ	3.258 (2)	H7C...C2	2.7800
C7...C6 ^x	3.451 (3)	H7C...H2	2.3300
C7...C4 ⁱⁱ	3.580 (3)	H7C...H6B ^x	2.5700
C1...H1 ⁱ	2.676 (15)	H7C...O2 ^v	2.5700
C1—O1—C5	121.64 (13)	C1—C2—H2	120.00
C3—O3—C7	117.64 (14)	C3—C2—H2	120.00
C6—O4—H1	108.3 (14)	C3—C4—H4	121.00
O1—C1—O2	114.29 (14)	C5—C4—H4	121.00
O1—C1—C2	117.44 (15)	O4—C6—H6A	110.00
O2—C1—C2	128.25 (16)	O4—C6—H6B	110.00
C1—C2—C3	120.23 (15)	C5—C6—H6A	110.00
O3—C3—C2	125.70 (15)	C5—C6—H6B	110.00
C2—C3—C4	120.35 (15)	H6A—C6—H6B	108.00
O3—C3—C4	113.95 (14)	O3—C7—H7A	109.00
C3—C4—C5	118.87 (15)	O3—C7—H7B	109.00
O1—C5—C4	121.46 (15)	O3—C7—H7C	109.00
C4—C5—C6	128.79 (16)	H7A—C7—H7B	109.00
O1—C5—C6	109.75 (14)	H7A—C7—H7C	109.00
O4—C6—C5	109.94 (14)	H7B—C7—H7C	109.00
C5—O1—C1—O2	179.27 (15)	C1—C2—C3—O3	178.79 (16)
C5—O1—C1—C2	0.3 (2)	C1—C2—C3—C4	-1.1 (3)
C1—O1—C5—C4	-0.4 (2)	O3—C3—C4—C5	-178.94 (15)
C1—O1—C5—C6	179.00 (15)	C2—C3—C4—C5	0.9 (2)
C7—O3—C3—C2	-0.9 (2)	C3—C4—C5—O1	-0.2 (2)
C7—O3—C3—C4	178.94 (15)	C3—C4—C5—C6	-179.49 (17)
O1—C1—C2—C3	0.4 (2)	O1—C5—C6—O4	177.77 (14)
O2—C1—C2—C3	-178.35 (18)	C4—C5—C6—O4	-2.9 (3)

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

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
O4—H1...O1 ⁱⁱⁱ	0.836 (14)	2.47 (2)	3.1840 (19)	144.2 (17)

O4—H1...O2 ⁱⁱⁱ	0.836 (14)	2.073 (13)	2.8678 (18)	158.6 (18)
C7—H7B...O4 ^{vi}	0.96	2.58	3.494 (2)	159
C7—H7C...O2 ^v	0.96	2.57	3.504 (2)	164

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