

2,3,4-Trihydroxybenzoic acid 0.25-hydrate

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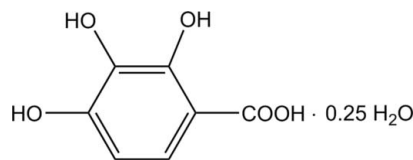
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Key indicators: single-crystal X-ray study; $T = 291$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; R factor = 0.036; wR factor = 0.090; data-to-parameter ratio = 11.5.

The asymmetric unit of the title compound, $\text{C}_7\text{H}_6\text{O}_5 \cdot 0.25\text{H}_2\text{O}$, contains two molecules of 2,3,4-trihydroxybenzoic acid, with similar conformations, and one water molecule which lies on a twofold rotation axis. Both acid molecules are essentially planar [maximum r.m.s deviations = 0.0324 (2) and 0.0542 (3) Å for the two acid molecules]. The molecular conformations are stabilized by intramolecular O(phenol)–H...O(carboxyl/phenol) interactions. A cyclic intermolecular association is formed between the two acid and one water molecule [graph set $R_3^3(12)$] involving O–H...O hydrogen bonds. The two acid molecules are further linked through a cyclic $R_2^2(8)$ carboxylic acid hydrogen-bonding association, which together with intermolecular O–H...O hydrogen-bonding interactions involving the phenol groups and the water molecule, and weak π – π interactions [minimum ring centroid separation = 3.731 (3) Å], give a three-dimensional network.

Related literature

For the natural distribution of 2,3,4-trihydroxybenzoic acid, see: Zhai *et al.* (2010); Xu & Chang (2010). For its antioxidant and antibacterial activities, see: Kodama *et al.* (2007); Friedman *et al.* (2003). For the inhibition of xanthine oxidase, see: Chang *et al.* (1995). For the crystal structure of the dihydrate pseudopolymorph, see: Prior & Sharp (2010). For π – π interactions in gallic acid pyridine monosolvate and in natural flavonoids, see: Dong *et al.* (2011); Jiang *et al.* (2009, 2002). For graph-set analysis, see: Etter *et al.* (1990); Bernstein *et al.* (1995).



Experimental

Crystal data

$\text{C}_7\text{H}_6\text{O}_5 \cdot 0.25\text{H}_2\text{O}$

$M_r = 174.62$

Orthorhombic, $P2_12_12_1$

$a = 11.8364$ (12) Å

$b = 32.598$ (3) Å

$c = 3.7306$ (4) Å

$V = 1439.4$ (3) Å³

$Z = 8$

Mo $K\alpha$ radiation

$\mu = 0.14$ mm⁻¹

$T = 291$ K

$0.42 \times 0.28 \times 0.20$ mm

Data collection

Bruker SMART 1000 CCD

diffractometer

Absorption correction: multi-scan

(*SADABS*; Sheldrick, 2004)

$T_{\min} = 0.874$, $T_{\max} = 1.000$

7863 measured reflections

2552 independent reflections

2010 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.060$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.036$

$wR(F^2) = 0.090$

$S = 0.97$

2552 reflections

222 parameters

H-atom parameters constrained

$\Delta\rho_{\text{max}} = 0.17$ e Å⁻³

$\Delta\rho_{\text{min}} = -0.15$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
$\text{O1W} - \text{H1WA} \cdots \text{O2}$	0.82	2.19	2.963 (2)	156
$\text{O1} - \text{H1A} \cdots \text{O4}$	0.82	1.88	2.587 (3)	143
$\text{O2} - \text{H2A} \cdots \text{O2}^{\text{i}}$	0.82	2.05	2.839 (4)	161
$\text{O3} - \text{H3A} \cdots \text{O3}^{\text{ii}}$	0.82	2.06	2.828 (3)	155
$\text{O5} - \text{H5B} \cdots \text{O4}^{\text{iii}}$	0.82	1.88	2.679 (4)	165
$\text{O1}' - \text{H1}'\text{A} \cdots \text{O4}'$	0.82	1.87	2.588 (3)	145
$\text{O2}' - \text{H2}'\text{A} \cdots \text{O1}$	0.82	1.95	2.729 (4)	159
$\text{O3}' - \text{H3}'\text{A} \cdots \text{O1W}$	0.82	2.06	2.841 (3)	158
$\text{O5}' - \text{H5}'\text{B} \cdots \text{O4}^{\text{iv}}$	0.82	1.85	2.659 (4)	171

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

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINT* (Bruker, 1998); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *XP* (Siemens, 1998); software used to prepare material for publication: *SHELXTL* (Sheldrick, 2008).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: ZS2181).

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supplementary materials

Acta Cryst. (2012). E68, o825–o826 [doi:10.1107/S160053681200709X]

2,3,4-Trihydroxybenzoic acid 0.25-hydrate**Jin-Hang Li, Fu-Yue Dong, Fang Cai, Xiao-Feng Yuan and Ren-Wang Jiang****Comment**

2,3,4-Trihydroxybenzoic acid is a polyphenolic acid which has been isolated from *Pachysandra terminalis* (Zhai *et al.*, 2010) and the lentil (Xu & Chang, 2010). It has been found to show antioxidant (Kodama *et al.*, 2007) and antibacterial activities (Friedman *et al.*, 2003) and also to inhibit the xanthine oxidase enzyme (Chang *et al.*, 1995). This compound contains two of the most common functional groups in natural products: the carboxylic acid and the phenolic groups. The dihydrate pseudopolymorph of the acid with a space group *P*-1 has previously been reported (Prior & Sharp, 2010). In this study, we report the structure of the second pseudopolymorph, the partial hydrate $C_7H_6O_5 \cdot 0.25H_2O$.

The asymmetric unit of the title compound contains two molecules of 2,3,4-trihydroxybenzoic acid [(1) and (2)] with similar conformations, and one water molecule of solvation which lies on a crystallographic twofold rotation axis (Fig. 1). Both acid molecules are essentially planar [maximum r.m.s deviation 0.0324 (2) and 0.0542 (3) Å for the two acid molecules], with a dihedral angle of 49.4 (3)° between the planes. The molecular conformation of the acid molecules is stabilized by a number of intramolecular phenolic O—H \cdots O interactions (Table 1).

The water molecule and the acid molecules are linked through O—H \cdots O hydrogen bonds (Table 1) with both acting as donors and acceptors, giving a cyclic association [graph set $R^3_3(12)$ (Etter *et al.*, 1990; Bernstein *et al.*, 1995)]. The 2,3,4-trihydroxybenzoic acid molecules form head-to-head pairs through intermolecular hydrogen bonds [graph set $R^2_2(8)$]. The molecular pairs are further extended into chains through hydrogen bond O2—H \cdots O2ⁱⁱ, while adjacent chains are connected *via* hydrogen bond O3—H \cdots O3ⁱⁱⁱ into a three-dimensional network (Fig. 2) [for symmetry codes (i) and (ii), see Table 1]. It is noteworthy that aromatic π - π associations also play an important role in the molecular packing. Adjacent 2,3,4-trihydroxybenzoic acid molecules form stacks which extend down the *c*-axis giving weak π - π interactions [minimum ring centroid separation, 3.731 (3) Å], which is larger than that in the dihydrate form of the acid (Prior & Sharp, 2010) and in gallic acid pyridine monosolvate (Dong *et al.*, 2011), but comparable with those in natural flavonoids (Jiang *et al.*, 2002, 2009).

Experimental

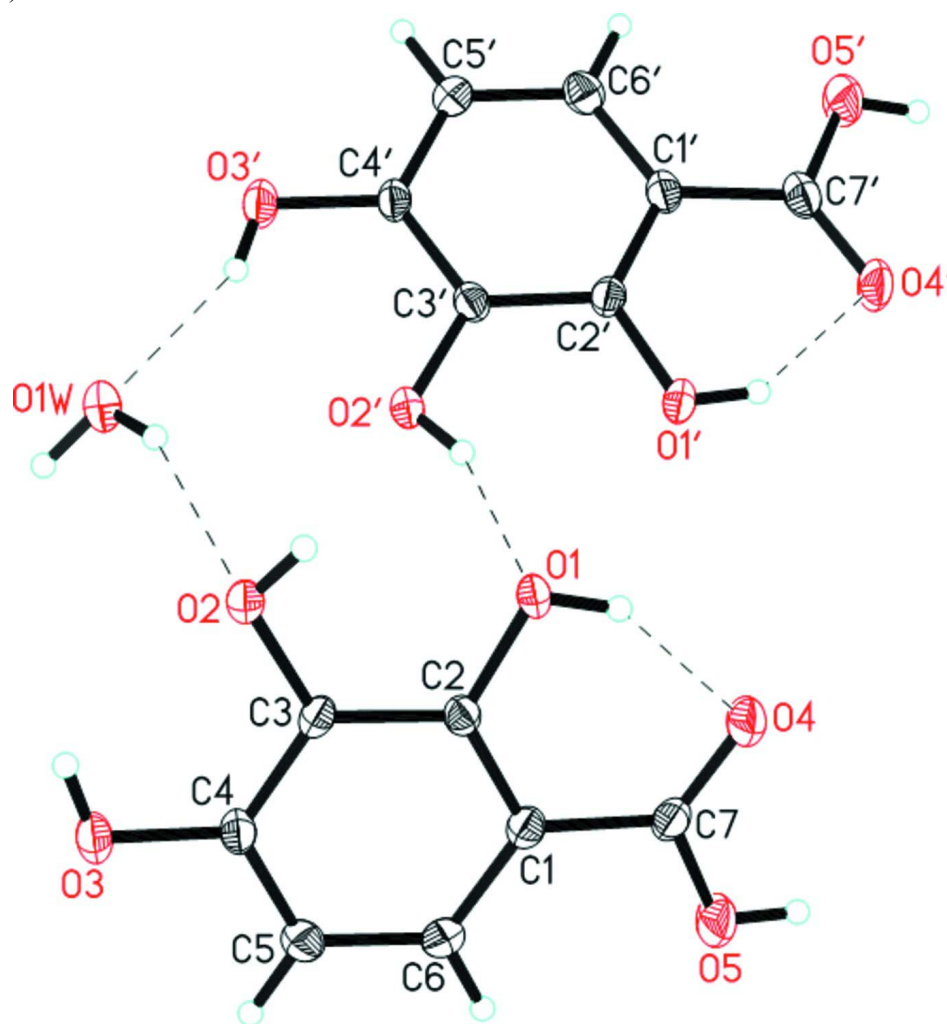
2,3,4-Trihydroxybenzoic acid (purity >98%) was purchased from the Sigma-Aldrich Company. Sample of 34 mg (0.2 mmol) was dissolved in methanol (1.5 ml) in a tube (4 ml) and sealed by parafilm. Light brown crystals were formed after three days at room temperature.

Refinement

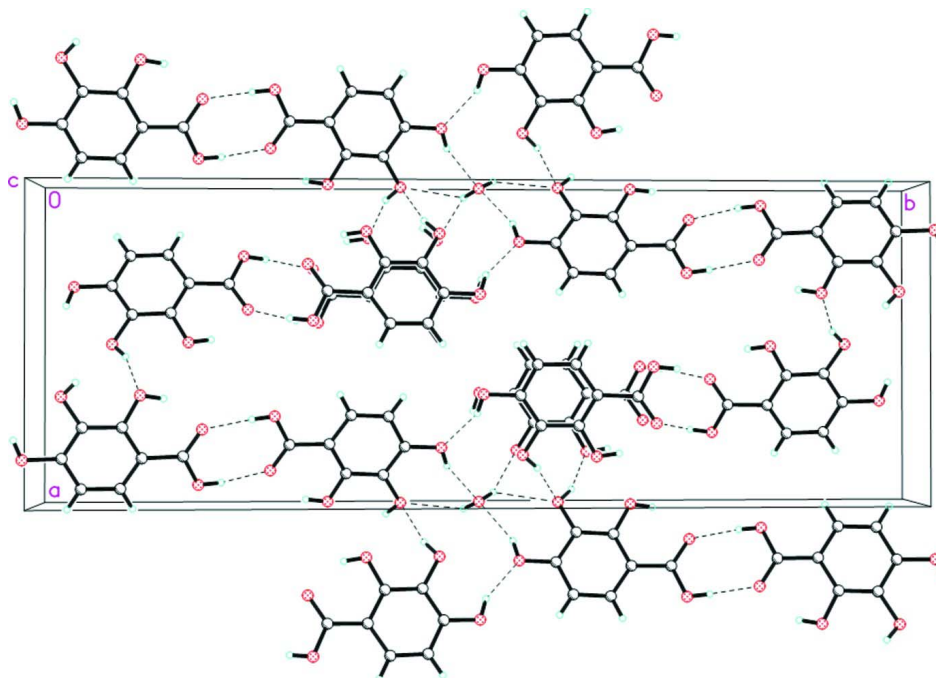
The C-bound H atoms were positioned geometrically and were included in the refinement in the riding-model approximation, with C—H = 0.96 Å (methyl C) and $U_{iso}(H) = 1.5U_{eq}(C)$; 0.97 Å (methylene C) and $U_{iso}(H) = 1.2U_{eq}(C)$; 0.93 Å (aryl H) and $U_{iso}(H) = 1.2U_{eq}(C)$; O—H = 0.82 Å and $U_{iso}(H) = 1.5U_{eq}(O)$.

Computing details

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINTE* (Bruker, 1998); data reduction: *SAINTE* (Bruker, 1998); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *XP* (Siemens, 1998); software used to prepare material for publication: *SHELXTL* (Sheldrick, 2008).

**Figure 1**

An ORTEP plot of the asymmetric unit of the title compound with atoms shown as 30% probability ellipsoids. The water molecule lies on a crystallographic twofold rotation axis and hydrogen bonds are shown as dashed lines.


Figure 2

Packing diagram viewed down the *c*-axis showing hydrogen bonds as dashed lines.

2,3,4-Trihydroxybenzoic acid 0.25-hydrate

Crystal data

$C_7H_6O_5 \cdot 0.25H_2O$

$M_r = 174.62$

Orthorhombic, $P2_12_12$

Hall symbol: $P\ 2ac\ 2ab$

$a = 11.8364\ (12)\ \text{\AA}$

$b = 32.598\ (3)\ \text{\AA}$

$c = 3.7306\ (4)\ \text{\AA}$

$V = 1439.4\ (3)\ \text{\AA}^3$

$Z = 8$

$F(000) = 724$

$D_x = 1.612\ \text{Mg m}^{-3}$

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

Cell parameters from 2552 reflections

$\theta = 2.1\text{--}50.0^\circ$

$\mu = 0.14\ \text{mm}^{-1}$

$T = 291\ \text{K}$

Prism, light brown

$0.42 \times 0.28 \times 0.20\ \text{mm}$

Data collection

Bruker SMART 1000 CCD
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

ω scan

Absorption correction: multi-scan
(*SADABS*; Sheldrick, 2004)

$T_{\min} = 0.874$, $T_{\max} = 1.000$

7863 measured reflections

2552 independent reflections

2010 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.060$

$\theta_{\max} = 25.0^\circ$, $\theta_{\min} = 2.1^\circ$

$h = -13 \rightarrow 14$

$k = -38 \rightarrow 34$

$l = -4 \rightarrow 4$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.036$

$wR(F^2) = 0.090$

$S = 0.97$

2552 reflections

222 parameters

0 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

$$w = 1/[\sigma^2(F_o^2) + (0.0443P)^2]$$

where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.17 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.15 \text{ e } \text{\AA}^{-3}$

Special details

Geometry. All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds 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 $F^2 > 2\text{sigma}(F^2)$ 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	$U_{\text{iso}}^*/U_{\text{eq}}$
O1W	0.5000	0.0000	-0.1102 (7)	0.0395 (6)
H1WA	0.5263	0.0173	0.0258	0.059*
O1	0.65345 (12)	0.12870 (4)	0.2193 (5)	0.0334 (4)
H1A	0.6637	0.1534	0.1900	0.040*
O2	0.65656 (13)	0.04688 (4)	0.3549 (4)	0.0323 (4)
H2A	0.6145	0.0640	0.4444	0.039*
O3	0.84576 (14)	0.00203 (5)	0.2010 (5)	0.0445 (5)
H3A	0.7859	-0.0048	0.2940	0.053*
O4	0.75826 (14)	0.19200 (4)	-0.0346 (6)	0.0429 (5)
O5	0.93058 (15)	0.18491 (4)	-0.2704 (6)	0.0481 (5)
H5B	0.9243	0.2098	-0.2919	0.058*
C1	0.84467 (19)	0.12653 (6)	-0.0250 (7)	0.0273 (5)
C2	0.74982 (19)	0.10787 (6)	0.1337 (6)	0.0245 (5)
C3	0.74932 (18)	0.06626 (6)	0.2074 (6)	0.0254 (5)
C4	0.84311 (19)	0.04271 (6)	0.1272 (7)	0.0295 (5)
C5	0.9374 (2)	0.06046 (7)	-0.0324 (7)	0.0330 (6)
H5A	0.9998	0.0444	-0.0891	0.040*
C6	0.9383 (2)	0.10180 (7)	-0.1064 (7)	0.0313 (6)
H6A	1.0018	0.1135	-0.2117	0.038*
C7	0.8406 (2)	0.17017 (7)	-0.1079 (7)	0.0310 (6)
O1'	0.49399 (13)	0.17376 (4)	-0.2824 (5)	0.0359 (4)
H1'A	0.4856	0.1982	-0.3270	0.043*
O2'	0.49999 (12)	0.09083 (4)	-0.2151 (5)	0.0308 (4)
H2'A	0.5380	0.1075	-0.1033	0.037*
O3'	0.32290 (14)	0.04439 (4)	-0.4474 (5)	0.0397 (5)
H3'A	0.3817	0.0378	-0.3464	0.048*
O4'	0.39072 (15)	0.23703 (5)	-0.5434 (6)	0.0446 (5)
O5'	0.22055 (15)	0.23003 (5)	-0.7935 (6)	0.0506 (5)
H5'B	0.2294	0.2547	-0.8259	0.061*
C1'	0.3088 (2)	0.17066 (7)	-0.5659 (7)	0.0293 (6)
C2'	0.40325 (19)	0.15242 (6)	-0.4052 (6)	0.0271 (6)
C3'	0.40863 (19)	0.11019 (6)	-0.3646 (6)	0.0267 (5)

C4'	0.31874 (19)	0.08593 (6)	-0.4795 (7)	0.0281 (6)
C5'	0.22322 (19)	0.10389 (7)	-0.6322 (7)	0.0328 (6)
H5'A	0.1630	0.0876	-0.7056	0.039*
C6'	0.2185 (2)	0.14569 (7)	-0.6737 (7)	0.0336 (6)
H6'A	0.1546	0.1576	-0.7746	0.040*
C7'	0.3098 (2)	0.21492 (7)	-0.6310 (7)	0.0349 (6)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1W	0.0469 (15)	0.0229 (11)	0.0486 (16)	-0.0056 (10)	0.000	0.000
O1	0.0299 (9)	0.0177 (7)	0.0527 (11)	0.0023 (6)	0.0057 (9)	0.0020 (8)
O2	0.0280 (9)	0.0212 (8)	0.0475 (11)	0.0003 (6)	0.0064 (8)	0.0020 (8)
O3	0.0386 (10)	0.0215 (8)	0.0734 (13)	0.0038 (7)	0.0121 (10)	0.0077 (10)
O4	0.0398 (11)	0.0244 (8)	0.0644 (13)	-0.0006 (8)	0.0084 (10)	0.0075 (9)
O5	0.0463 (11)	0.0268 (9)	0.0713 (14)	-0.0031 (8)	0.0157 (11)	0.0094 (10)
C1	0.0298 (13)	0.0244 (11)	0.0278 (13)	-0.0029 (10)	-0.0040 (11)	0.0009 (11)
C2	0.0265 (12)	0.0208 (11)	0.0261 (13)	0.0014 (9)	-0.0058 (11)	-0.0024 (10)
C3	0.0285 (13)	0.0211 (11)	0.0268 (12)	-0.0056 (9)	-0.0017 (11)	0.0004 (10)
C4	0.0334 (14)	0.0193 (11)	0.0359 (14)	0.0014 (10)	-0.0033 (12)	0.0009 (11)
C5	0.0278 (14)	0.0303 (13)	0.0409 (15)	0.0041 (10)	0.0006 (12)	-0.0034 (12)
C6	0.0257 (13)	0.0324 (13)	0.0360 (15)	-0.0042 (10)	-0.0001 (12)	0.0006 (11)
C7	0.0315 (14)	0.0255 (12)	0.0361 (14)	-0.0068 (11)	-0.0040 (12)	0.0019 (11)
O1'	0.0346 (9)	0.0185 (7)	0.0546 (11)	-0.0032 (6)	-0.0058 (10)	0.0011 (8)
O2'	0.0294 (9)	0.0191 (7)	0.0440 (10)	0.0016 (6)	-0.0038 (9)	0.0011 (7)
O3'	0.0408 (11)	0.0207 (8)	0.0575 (12)	-0.0029 (7)	-0.0115 (9)	-0.0004 (9)
O4'	0.0476 (12)	0.0208 (8)	0.0655 (14)	0.0004 (8)	-0.0142 (10)	0.0057 (9)
O5'	0.0482 (11)	0.0279 (9)	0.0758 (14)	0.0047 (8)	-0.0172 (12)	0.0138 (10)
C1'	0.0340 (14)	0.0227 (12)	0.0312 (13)	0.0031 (10)	0.0040 (11)	-0.0003 (11)
C2'	0.0295 (13)	0.0221 (12)	0.0297 (14)	-0.0024 (10)	0.0035 (12)	-0.0021 (10)
C3'	0.0280 (13)	0.0231 (11)	0.0290 (13)	0.0037 (10)	0.0045 (11)	0.0009 (10)
C4'	0.0334 (14)	0.0199 (11)	0.0311 (14)	0.0005 (10)	0.0052 (11)	-0.0007 (11)
C5'	0.0297 (14)	0.0299 (12)	0.0388 (15)	-0.0033 (10)	-0.0005 (12)	-0.0019 (12)
C6'	0.0342 (14)	0.0326 (12)	0.0339 (15)	0.0051 (10)	-0.0039 (12)	0.0022 (12)
C7'	0.0399 (16)	0.0271 (13)	0.0376 (15)	0.0071 (11)	0.0020 (12)	0.0021 (12)

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

O1W—H1WA	0.8200	O1'—C2'	1.359 (3)
O1—C2	1.365 (3)	O1'—H1'A	0.8200
O1—H1A	0.8200	O2'—C3'	1.371 (3)
O2—C3	1.381 (3)	O2'—H2'A	0.8200
O2—H2A	0.8200	O3'—C4'	1.360 (2)
O3—C4	1.355 (2)	O3'—H3'A	0.8200
O3—H3A	0.8200	O4'—C7'	1.243 (3)
O4—C7	1.237 (3)	O5'—C7'	1.314 (3)
O5—C7	1.317 (3)	O5'—H5'B	0.8200
O5—H5B	0.8200	C1'—C2'	1.401 (3)
C1—C6	1.404 (3)	C1'—C6'	1.403 (3)
C1—C2	1.407 (3)	C1'—C7'	1.463 (3)

C1—C7	1.457 (3)	C2'—C3'	1.386 (3)
C2—C3	1.384 (3)	C3'—C4'	1.393 (3)
C3—C4	1.382 (3)	C4'—C5'	1.395 (3)
C4—C5	1.391 (3)	C5'—C6'	1.373 (3)
C5—C6	1.376 (3)	C5'—H5'A	0.9300
C5—H5A	0.9300	C6'—H6'A	0.9300
C6—H6A	0.9300		
C2—O1—H1A	109.5	C2'—O1'—H1'A	109.5
C3—O2—H2A	109.5	C3'—O2'—H2'A	109.5
C4—O3—H3A	109.5	C4'—O3'—H3'A	109.5
C7—O5—H5B	109.5	C7'—O5'—H5'B	109.5
C6—C1—C2	118.2 (2)	C2'—C1'—C6'	119.0 (2)
C6—C1—C7	122.8 (2)	C2'—C1'—C7'	118.9 (2)
C2—C1—C7	119.0 (2)	C6'—C1'—C7'	122.1 (2)
O1—C2—C3	115.9 (2)	O1'—C2'—C3'	115.8 (2)
O1—C2—C1	123.35 (18)	O1'—C2'—C1'	123.89 (18)
C3—C2—C1	120.7 (2)	C3'—C2'—C1'	120.3 (2)
O2—C3—C4	118.08 (17)	O2'—C3'—C2'	122.5 (2)
O2—C3—C2	122.06 (19)	O2'—C3'—C4'	117.79 (18)
C4—C3—C2	119.9 (2)	C2'—C3'—C4'	119.7 (2)
O3—C4—C3	121.2 (2)	O3'—C4'—C3'	120.7 (2)
O3—C4—C5	118.4 (2)	O3'—C4'—C5'	118.9 (2)
C3—C4—C5	120.4 (2)	C3'—C4'—C5'	120.4 (2)
C6—C5—C4	119.9 (2)	C6'—C5'—C4'	119.7 (2)
C6—C5—H5A	120.0	C6'—C5'—H5'A	120.1
C4—C5—H5A	120.0	C4'—C5'—H5'A	120.1
C5—C6—C1	120.9 (2)	C5'—C6'—C1'	120.8 (2)
C5—C6—H6A	119.6	C5'—C6'—H6'A	119.6
C1—C6—H6A	119.6	C1'—C6'—H6'A	119.6
O4—C7—O5	122.0 (2)	O4'—C7'—O5'	121.6 (2)
O4—C7—C1	122.7 (2)	O4'—C7'—C1'	122.3 (2)
O5—C7—C1	115.3 (2)	O5'—C7'—C1'	116.1 (2)
C6—C1—C2—O1	-178.9 (2)	C6'—C1'—C2'—O1'	-178.2 (2)
C7—C1—C2—O1	-0.8 (3)	C7'—C1'—C2'—O1'	4.6 (4)
C6—C1—C2—C3	0.1 (3)	C6'—C1'—C2'—C3'	2.1 (4)
C7—C1—C2—C3	178.2 (2)	C7'—C1'—C2'—C3'	-175.1 (2)
O1—C2—C3—O2	0.7 (3)	O1'—C2'—C3'—O2'	-0.9 (3)
C1—C2—C3—O2	-178.4 (2)	C1'—C2'—C3'—O2'	178.8 (2)
O1—C2—C3—C4	179.6 (2)	O1'—C2'—C3'—C4'	179.3 (2)
C1—C2—C3—C4	0.5 (3)	C1'—C2'—C3'—C4'	-1.0 (4)
O2—C3—C4—O3	-2.0 (3)	O2'—C3'—C4'—O3'	-0.8 (3)
C2—C3—C4—O3	179.0 (2)	C2'—C3'—C4'—O3'	179.0 (2)
O2—C3—C4—C5	177.9 (2)	O2'—C3'—C4'—C5'	179.7 (2)
C2—C3—C4—C5	-1.0 (4)	C2'—C3'—C4'—C5'	-0.5 (4)
O3—C4—C5—C6	-179.1 (2)	O3'—C4'—C5'—C6'	-178.6 (2)
C3—C4—C5—C6	1.0 (4)	C3'—C4'—C5'—C6'	0.9 (4)
C4—C5—C6—C1	-0.4 (4)	C4'—C5'—C6'—C1'	0.3 (4)

C2—C1—C6—C5	-0.1 (4)	C2'—C1'—C6'—C5'	-1.8 (4)
C7—C1—C6—C5	-178.2 (2)	C7'—C1'—C6'—C5'	175.3 (2)
C6—C1—C7—O4	-179.6 (2)	C2'—C1'—C7'—O4'	-1.2 (4)
C2—C1—C7—O4	2.3 (4)	C6'—C1'—C7'—O4'	-178.3 (2)
C6—C1—C7—O5	1.0 (3)	C2'—C1'—C7'—O5'	177.7 (2)
C2—C1—C7—O5	-177.1 (2)	C6'—C1'—C7'—O5'	0.6 (4)

Hydrogen-bond geometry (Å, °)

<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>
O1 <i>W</i> —H1 <i>WA</i> ...O2	0.82	2.19	2.963 (2)	156
O1 <i>W</i> —H1 <i>WA</i> ...O2'	0.82	2.58	2.987 (3)	112
O1—H1 <i>A</i> ...O4	0.82	1.88	2.587 (3)	143
O2—H2 <i>A</i> ...O2 ⁱⁱ	0.82	2.05	2.839 (4)	161
O2—H2 <i>A</i> ...O1	0.82	2.32	2.715 (2)	111
O3—H3 <i>A</i> ...O3 ⁱⁱⁱ	0.82	2.06	2.828 (3)	155
O3—H3 <i>A</i> ...O2	0.82	2.29	2.735 (3)	115
O5—H5 <i>B</i> ...O4 ⁱⁱⁱ	0.82	1.88	2.679 (4)	165
O1'—H1' <i>A</i> ...O4'	0.82	1.87	2.588 (3)	145
O2'—H2' <i>A</i> ...O1	0.82	1.95	2.729 (4)	159
O2'—H2' <i>A</i> ...O1'	0.82	2.32	2.716 (2)	110
O3'—H3' <i>A</i> ...O1 <i>W</i>	0.82	2.06	2.841 (3)	158
O3'—H3' <i>A</i> ...O2'	0.82	2.28	2.727 (4)	115
O5'—H5' <i>B</i> ...O4 ^{iv}	0.82	1.85	2.659 (4)	171

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