

Biphenyl-2,4,4',6-tetracarboxylic acid monohydrate

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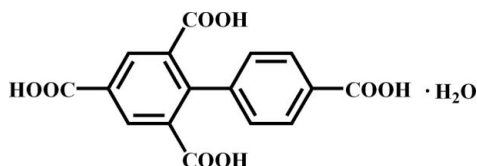
Received 11 July 2013; accepted 16 July 2013

Key indicators: single-crystal X-ray study; $T = 296$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; R factor = 0.052; wR factor = 0.152; data-to-parameter ratio = 15.2.

In the title compound, $\text{C}_{16}\text{H}_{10}\text{O}_8 \cdot \text{H}_2\text{O}$, the dihedral angle between the benzene rings is $71.59(8)^\circ$. The COOH groups make dihedral angles of $10.3(2)$, $30.8(2)$, $11.3(2)$ and $42.3(2)^\circ$ with their attached rings. In the crystal, $\text{O}-\text{H} \cdots \text{O}$ hydrogen bonds link the components forming a three-dimensional supramolecular network.

Related literature

For general background to the use of aromatic carboxylates as building blocks for the construction of various architectures, see: Yaghi *et al.* (2003); Zhao *et al.* (2012).



Experimental

Crystal data

$\text{C}_{16}\text{H}_{10}\text{O}_8 \cdot \text{H}_2\text{O}$
 $M_r = 348.26$
Monoclinic, $P2_1/c$
 $a = 5.638(4)$ Å
 $b = 16.160(11)$ Å
 $c = 16.798(12)$ Å
 $\beta = 92.524(12)^\circ$

$V = 1528.9(19)$ Å³
 $Z = 4$
Mo $K\alpha$ radiation
 $\mu = 0.13$ mm⁻¹
 $T = 296$ K
 $0.21 \times 0.18 \times 0.17$ mm

Data collection

Bruker SMART CCD diffractometer
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.974$, $T_{\max} = 0.979$

15936 measured reflections
3516 independent reflections
3070 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.124$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.052$
 $wR(F^2) = 0.152$
 $S = 1.02$
3516 reflections
232 parameters
3 restraints

H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\text{max}} = 0.30$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.26$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H} \cdots A$	$D-\text{H}$	$\text{H} \cdots A$	$D \cdots A$	$D-\text{H} \cdots A$
$\text{O2}-\text{H2A} \cdots \text{O3}^{\text{i}}$	0.82	1.87	2.688 (2)	179
$\text{O4}-\text{H4B} \cdots \text{O1W}^{\text{ii}}$	0.82	1.91	2.725 (2)	177
$\text{O6}-\text{H6B} \cdots \text{O8}^{\text{iii}}$	0.82	1.82	2.636 (2)	178
$\text{O7}-\text{H7B} \cdots \text{O1W}^{\text{iv}}$	0.82	1.87	2.688 (2)	176
$\text{O1W}-\text{H1WB} \cdots \text{O1}$	0.85 (1)	2.09 (1)	2.818 (2)	143 (2)
$\text{O1W}-\text{H1WA} \cdots \text{O5}^{\text{v}}$	0.86 (1)	1.98 (1)	2.787 (2)	157 (2)

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

Data collection: SMART (Bruker, 2007); cell refinement: SAINT (Bruker, 2007); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: SHELXTL (Sheldrick, 2008); software used to prepare material for publication: SHELXTL.

The authors are grateful to the Natural Science Foundation of Hubei Province of China (grant No. 2010CDB10707) and the Project of Hubei Provincial Education Office (grant No. Q20101203).

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: LH5634).

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Zhao, J., Li, D. S., Ke, X. J., Liu, B., Zou, K. & Hu, H. M. (2012). Dalton Trans. 41, 2560–2563.

supporting information

Acta Cryst. (2013). E69, o1373 [doi:10.1107/S1600536813019624]

Biphenyl-2,4,4',6-tetracarboxylic acid monohydrate**Ye-Nan Wang and Jun Zhao****S1. Comment**

Aromatic carboxylates have been proven to be effective building blocks for the design and construction of coordination polymers exhibiting remarkable polymeric structural motifs due to their rich coordination modes (Yaghi *et al.*, 2003; Zhao *et al.*, 2012). Recently, we attempted to synthesize an Sm^{III} complex with the ligand in hydrothermal synthesis conditions. However the title organic salt was obtained, its structure is reported here.

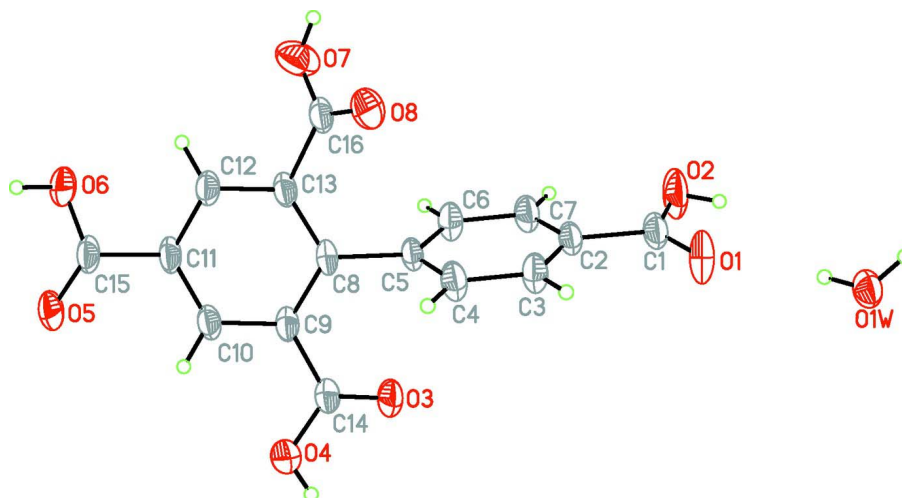
The molecular structure of the title compound is shown in Fig. 1. The asymmetric unit comprises one biphenyl-1,2,4,4',6-tetracarboxylic acid and one solvent water molecule. The dihedral angle between the two benzene rings of the biphenyl-1,2,4,4',6-tetracarboxylic acid is 71.59 (8)°. This may be the result of intermolecular O—H...O interactions and steric effects. In the crystal, extensive O—H...O hydrogen bonds involving biphenyl-1,2,4,4',6-tetracarboxylic acid molecules and water molecules link the components into a three-dimensional supramolecular network (Fig. 2).

S2. Experimental

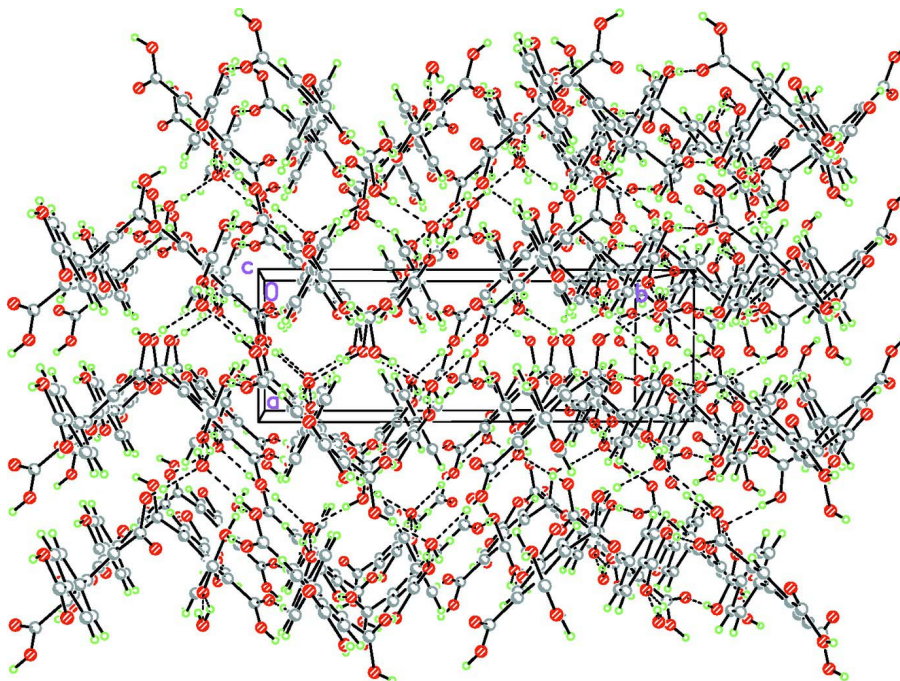
All chemicals were of reagent grade quality obtained from commercial sources and used without further purification. A mixture of biphenyl-1,2,4,4',6-tetracarboxylic acid (0.0370 g, 0.1 mmol), Sm(NO₃)₃·6H₂O (0.0444 g, 0.1 mmol) and water (12 ml) were placed in a 23 ml Teflon-lined stainless steel reactor and heated at 393 K for 2 days, and then cooled to room temperature at 10 K h⁻¹ to obtain colorless prism-shaped crystals suitable for X-ray analysis.

S3. Refinement

The H atoms bonded to C and carboxylate O atoms were positioned geometrically (C—H = 0.93, O—H = 0.82 Å) and allowed to ride on their parent atoms, with $U_{\text{iso}}(\text{H})$ value equal to $1.2U_{\text{eq}}(\text{C})$ or $1.5U_{\text{eq}}(\text{O})$. The H atoms bonded to water were located in a difference Fourier map and refined with O—H distance restraint of 0.85 ± 0.01 Å, $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{O})$.

**Figure 1**

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

**Figure 2**

Part of the crystal structure with hydrogen bonds shown as dashed lines.

Biphenyl-2,4,4',6-tetracarboxylic acid monohydrate

Crystal data

$C_{16}H_{10}O_8 \cdot H_2O$

$M_r = 348.26$

Monoclinic, $P2_1/c$

Hall symbol: $-p\ 2ybc$

$a = 5.638\ (4)\ \text{\AA}$

$b = 16.160\ (11)\ \text{\AA}$

$c = 16.798\ (12)\ \text{\AA}$

$\beta = 92.524\ (12)^\circ$

$V = 1528.9\ (19)\ \text{\AA}^3$

$Z = 4$

$F(000) = 720$

$D_x = 1.513\ \text{Mg m}^{-3}$

Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$
 Cell parameters from 4298 reflections
 $\theta = 3.5\text{--}27.6^\circ$
 $\mu = 0.13 \text{ mm}^{-1}$

$T = 296 \text{ K}$
 Prism, colorless
 $0.21 \times 0.18 \times 0.17 \text{ mm}$

Data collection

Bruker SMART CCD
 diffractometer
 Radiation source: fine-focus sealed tube
 Graphite monochromator
 φ and ω scans
 Absorption correction: multi-scan
 (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.974$, $T_{\max} = 0.979$

15936 measured reflections
 3516 independent reflections
 3070 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.124$
 $\theta_{\max} = 27.6^\circ$, $\theta_{\min} = 3.5^\circ$
 $h = -7 \rightarrow 7$
 $k = -20 \rightarrow 20$
 $l = -21 \rightarrow 21$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.052$
 $wR(F^2) = 0.152$
 $S = 1.02$
 3516 reflections
 232 parameters
 3 restraints
 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
 $w = 1/[\sigma^2(F_o^2) + (0.0821P)^2 + 0.3015P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.30 \text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.26 \text{ e \AA}^{-3}$

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	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	-0.0695 (3)	0.89761 (9)	0.06051 (8)	0.0368 (3)
C2	-0.0558 (3)	0.89091 (9)	-0.02802 (8)	0.0319 (3)
C3	0.1413 (3)	0.85407 (10)	-0.06078 (8)	0.0369 (3)
H3A	0.2633	0.8333	-0.0274	0.044*
C4	0.1565 (3)	0.84821 (10)	-0.14271 (8)	0.0368 (3)
H4A	0.2885	0.8236	-0.1641	0.044*
C5	-0.0247 (2)	0.87892 (8)	-0.19285 (7)	0.0298 (3)
C6	-0.2205 (3)	0.91575 (10)	-0.15985 (8)	0.0367 (3)
H6A	-0.3424	0.9365	-0.1932	0.044*
C7	-0.2365 (3)	0.92190 (10)	-0.07773 (8)	0.0366 (3)
H7A	-0.3682	0.9468	-0.0563	0.044*
C8	-0.0256 (2)	0.87106 (8)	-0.28226 (7)	0.0291 (3)

C9	0.1157 (2)	0.92037 (8)	-0.33066 (7)	0.0303 (3)
C10	0.1001 (3)	0.91203 (9)	-0.41364 (8)	0.0336 (3)
H10A	0.2018	0.9420	-0.4448	0.040*
C11	-0.0664 (3)	0.85925 (9)	-0.44958 (7)	0.0336 (3)
C12	-0.2109 (3)	0.81166 (9)	-0.40302 (8)	0.0332 (3)
H12A	-0.3254	0.7773	-0.4270	0.040*
C13	-0.1844 (2)	0.81537 (8)	-0.32018 (7)	0.0305 (3)
C14	0.2767 (2)	0.98523 (9)	-0.29572 (8)	0.0316 (3)
C15	-0.0870 (3)	0.85389 (10)	-0.53843 (8)	0.0375 (3)
C16	-0.3229 (3)	0.75439 (9)	-0.27373 (8)	0.0325 (3)
O1W	0.24574 (18)	0.88370 (7)	0.26856 (6)	0.0378 (3)
H1WB	0.159 (3)	0.8987 (13)	0.2288 (8)	0.057*
O1	0.0881 (3)	0.87787 (9)	0.10711 (6)	0.0544 (4)
O2	-0.2753 (2)	0.92774 (10)	0.08217 (7)	0.0603 (4)
H2A	-0.2625	0.9434	0.1286	0.090*
O3	0.2393 (2)	1.02125 (8)	-0.23453 (6)	0.0467 (3)
O4	0.4623 (2)	1.00041 (8)	-0.33852 (7)	0.0479 (3)
H4B	0.5466	1.0353	-0.3159	0.072*
O5	0.0525 (3)	0.88594 (9)	-0.58157 (6)	0.0556 (4)
O6	-0.2751 (3)	0.81178 (9)	-0.56371 (6)	0.0564 (4)
H6B	-0.2663	0.8019	-0.6114	0.085*
O7	-0.5391 (2)	0.74324 (9)	-0.30234 (8)	0.0545 (4)
H7B	-0.5991	0.7040	-0.2796	0.082*
O8	-0.2375 (2)	0.71750 (8)	-0.21700 (7)	0.0511 (3)
H1WA	0.158 (3)	0.8750 (16)	0.3080 (9)	0.077*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0543 (8)	0.0375 (7)	0.0190 (6)	-0.0055 (6)	0.0060 (5)	-0.0034 (5)
C2	0.0443 (7)	0.0347 (7)	0.0171 (6)	-0.0052 (6)	0.0058 (5)	-0.0024 (5)
C3	0.0435 (7)	0.0483 (8)	0.0189 (6)	0.0039 (6)	0.0000 (5)	-0.0002 (5)
C4	0.0403 (7)	0.0500 (8)	0.0205 (6)	0.0055 (6)	0.0056 (5)	-0.0026 (6)
C5	0.0403 (7)	0.0347 (7)	0.0147 (6)	-0.0033 (5)	0.0041 (5)	-0.0009 (5)
C6	0.0408 (7)	0.0492 (8)	0.0201 (6)	0.0070 (6)	0.0010 (5)	-0.0016 (5)
C7	0.0421 (7)	0.0463 (8)	0.0219 (6)	0.0034 (6)	0.0065 (5)	-0.0053 (5)
C8	0.0387 (6)	0.0347 (7)	0.0142 (5)	0.0040 (5)	0.0038 (5)	-0.0009 (5)
C9	0.0413 (7)	0.0324 (7)	0.0173 (6)	0.0012 (5)	0.0031 (5)	-0.0014 (5)
C10	0.0469 (7)	0.0364 (7)	0.0179 (6)	-0.0033 (6)	0.0069 (5)	-0.0003 (5)
C11	0.0519 (8)	0.0349 (7)	0.0144 (6)	-0.0012 (6)	0.0042 (5)	-0.0004 (5)
C12	0.0479 (7)	0.0346 (7)	0.0173 (6)	-0.0039 (6)	0.0019 (5)	-0.0018 (5)
C13	0.0423 (7)	0.0328 (7)	0.0167 (6)	0.0003 (5)	0.0048 (5)	0.0009 (5)
C14	0.0396 (7)	0.0357 (7)	0.0196 (6)	0.0014 (5)	0.0019 (5)	0.0008 (5)
C15	0.0583 (9)	0.0386 (8)	0.0157 (6)	-0.0041 (6)	0.0040 (5)	-0.0024 (5)
C16	0.0468 (7)	0.0336 (7)	0.0177 (6)	-0.0005 (6)	0.0059 (5)	-0.0008 (5)
O1W	0.0384 (5)	0.0507 (6)	0.0245 (5)	0.0005 (4)	0.0056 (4)	-0.0009 (4)
O1	0.0762 (8)	0.0685 (9)	0.0182 (5)	0.0146 (7)	-0.0019 (5)	-0.0023 (5)
O2	0.0637 (7)	0.0971 (11)	0.0207 (5)	0.0110 (7)	0.0100 (5)	-0.0130 (6)

O3	0.0637 (7)	0.0513 (7)	0.0259 (5)	-0.0128 (5)	0.0106 (5)	-0.0143 (4)
O4	0.0489 (6)	0.0598 (7)	0.0362 (6)	-0.0160 (5)	0.0131 (5)	-0.0146 (5)
O5	0.0808 (9)	0.0689 (8)	0.0179 (5)	-0.0253 (7)	0.0126 (5)	-0.0035 (5)
O6	0.0769 (9)	0.0744 (9)	0.0178 (5)	-0.0256 (7)	0.0019 (5)	-0.0063 (5)
O7	0.0483 (6)	0.0621 (8)	0.0528 (8)	-0.0122 (6)	-0.0002 (5)	0.0241 (6)
O8	0.0701 (8)	0.0563 (7)	0.0265 (5)	-0.0137 (6)	-0.0031 (5)	0.0156 (5)

Geometric parameters (Å, °)

C1—O1	1.201 (2)	C10—H10A	0.9300
C1—O2	1.324 (2)	C11—C12	1.387 (2)
C1—C2	1.496 (2)	C11—C15	1.494 (2)
C2—C7	1.382 (2)	C12—C13	1.394 (2)
C2—C3	1.395 (2)	C12—H12A	0.9300
C3—C4	1.386 (2)	C13—C16	1.498 (2)
C3—H3A	0.9300	C14—O3	1.2079 (19)
C4—C5	1.387 (2)	C14—O4	1.3181 (19)
C4—H4A	0.9300	C15—O5	1.210 (2)
C5—C6	1.391 (2)	C15—O6	1.314 (2)
C5—C8	1.507 (2)	C16—O8	1.2061 (19)
C6—C7	1.390 (2)	C16—O7	1.303 (2)
C6—H6A	0.9300	O1W—H1WB	0.845 (9)
C7—H7A	0.9300	O1W—H1WA	0.856 (9)
C8—C13	1.403 (2)	O2—H2A	0.8200
C8—C9	1.4099 (19)	O4—H4B	0.8200
C9—C10	1.399 (2)	O6—H6B	0.8200
C9—C14	1.490 (2)	O7—H7B	0.8200
C10—C11	1.387 (2)		
O1—C1—O2	123.42 (14)	C11—C10—C9	120.31 (13)
O1—C1—C2	124.02 (15)	C11—C10—H10A	119.8
O2—C1—C2	112.56 (13)	C9—C10—H10A	119.8
C7—C2—C3	119.67 (13)	C12—C11—C10	119.92 (13)
C7—C2—C1	120.35 (13)	C12—C11—C15	120.62 (13)
C3—C2—C1	119.97 (13)	C10—C11—C15	119.46 (13)
C4—C3—C2	120.40 (13)	C11—C12—C13	119.94 (13)
C4—C3—H3A	119.8	C11—C12—H12A	120.0
C2—C3—H3A	119.8	C13—C12—H12A	120.0
C3—C4—C5	120.16 (13)	C12—C13—C8	121.27 (12)
C3—C4—H4A	119.9	C12—C13—C16	117.14 (12)
C5—C4—H4A	119.9	C8—C13—C16	121.51 (12)
C4—C5—C6	119.18 (13)	O3—C14—O4	123.16 (14)
C4—C5—C8	123.02 (12)	O3—C14—C9	123.23 (13)
C6—C5—C8	117.74 (12)	O4—C14—C9	113.60 (12)
C7—C6—C5	120.91 (13)	O5—C15—O6	124.40 (14)
C7—C6—H6A	119.5	O5—C15—C11	123.34 (14)
C5—C6—H6A	119.5	O6—C15—C11	112.26 (13)
C2—C7—C6	119.68 (13)	O8—C16—O7	123.88 (14)

C2—C7—H7A	120.2	O8—C16—C13	122.53 (14)
C6—C7—H7A	120.2	O7—C16—C13	113.54 (12)
C13—C8—C9	117.81 (12)	H1WB—O1W—H1WA	108.9 (14)
C13—C8—C5	118.72 (12)	C1—O2—H2A	109.5
C9—C8—C5	123.37 (12)	C14—O4—H4B	109.5
C10—C9—C8	120.50 (13)	C15—O6—H6B	109.5
C10—C9—C14	118.09 (12)	C16—O7—H7B	109.5
C8—C9—C14	121.35 (12)		
O1—C1—C2—C7	-174.12 (16)	C14—C9—C10—C11	-172.69 (13)
O2—C1—C2—C7	5.9 (2)	C9—C10—C11—C12	-2.9 (2)
O1—C1—C2—C3	5.1 (2)	C9—C10—C11—C15	177.47 (13)
O2—C1—C2—C3	-174.80 (15)	C10—C11—C12—C13	-1.6 (2)
C7—C2—C3—C4	-0.2 (2)	C15—C11—C12—C13	177.95 (13)
C1—C2—C3—C4	-179.45 (14)	C11—C12—C13—C8	4.8 (2)
C2—C3—C4—C5	-0.1 (2)	C11—C12—C13—C16	-172.06 (13)
C3—C4—C5—C6	0.2 (2)	C9—C8—C13—C12	-3.2 (2)
C3—C4—C5—C8	-177.01 (14)	C5—C8—C13—C12	173.20 (13)
C4—C5—C6—C7	-0.1 (2)	C9—C8—C13—C16	173.50 (12)
C8—C5—C6—C7	177.24 (13)	C5—C8—C13—C16	-10.09 (19)
C3—C2—C7—C6	0.3 (2)	C10—C9—C14—O3	148.67 (15)
C1—C2—C7—C6	179.54 (14)	C8—C9—C14—O3	-28.5 (2)
C5—C6—C7—C2	-0.1 (2)	C10—C9—C14—O4	-30.44 (18)
C4—C5—C8—C13	107.63 (17)	C8—C9—C14—O4	152.42 (13)
C6—C5—C8—C13	-69.63 (18)	C12—C11—C15—O5	-170.64 (17)
C4—C5—C8—C9	-76.17 (19)	C10—C11—C15—O5	9.0 (2)
C6—C5—C8—C9	106.56 (17)	C12—C11—C15—O6	10.4 (2)
C13—C8—C9—C10	-1.40 (19)	C10—C11—C15—O6	-170.01 (15)
C5—C8—C9—C10	-177.63 (12)	C12—C13—C16—O8	136.42 (16)
C13—C8—C9—C14	175.67 (12)	C8—C13—C16—O8	-40.4 (2)
C5—C8—C9—C14	-0.6 (2)	C12—C13—C16—O7	-41.20 (19)
C8—C9—C10—C11	4.5 (2)	C8—C13—C16—O7	141.96 (15)

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>
O2—H2A...O3 ⁱ	0.82	1.87	2.688 (2)	179
O4—H4B...O1W ⁱⁱ	0.82	1.91	2.725 (2)	177
O6—H6B...O8 ⁱⁱⁱ	0.82	1.82	2.636 (2)	178
O7—H7B...O1W ^{iv}	0.82	1.87	2.688 (2)	176
O1W—H1WB...O1	0.85 (1)	2.09 (1)	2.818 (2)	143 (2)
O1W—H1WA...O5 ^v	0.86 (1)	1.98 (1)	2.787 (2)	157 (2)

Symmetry codes: (i) $-x, -y+2, -z$; (ii) $-x+1, -y+2, -z$; (iii) $x, -y+3/2, z-1/2$; (iv) $x-1, -y+3/2, z-1/2$; (v) $x, y, z+1$.