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(1*S*,2*S*,6*S*,9*S*)-6-Methyl-5-oxobicyclo-[4.4.0]decane-2,9-diol diacetate

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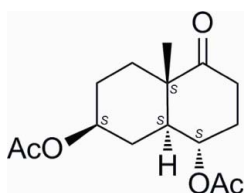
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Key indicators: single-crystal X-ray study; $T = 300$ K; mean $\sigma(\text{C}-\text{C}) = 0.004$ Å; R factor = 0.037; wR factor = 0.099; data-to-parameter ratio = 7.6.

The chiral title compound, $\text{C}_{15}\text{H}_{22}\text{O}_5$, is an intermediate in the total synthesis of biologically active 9,11-secoosterols. In the crystal, the cyclohexane rings are *trans*-fused and both adopt chair conformations. In the crystal, molecules are loosely held together in a layer parallel to (100) by weak intermolecular C—H...O hydrogen bonds accepted by carbonyl O atoms of the acetyl groups.

Related literature

For background to the biological activity of 9,11-secoosterols and the synthesis of the title compound, see: Aav *et al.* (2000). For a related structure, see: Foot *et al.* (2006). For hydrogen bonding, see: Steiner (2002).



Experimental

Crystal data

$\text{C}_{15}\text{H}_{22}\text{O}_5$
 $M_r = 282.33$
 Monoclinic, C_2
 $a = 22.885$ (5) Å
 $b = 9.340$ (2) Å

$c = 7.2250$ (13) Å
 $\beta = 101.280$ (6)°
 $V = 1514.5$ (5) Å³
 $Z = 4$
 Mo $K\alpha$ radiation

$\mu = 0.09$ mm⁻¹
 $T = 300$ K

0.50 × 0.20 × 0.16 mm

Data collection

Bruker SMART X2S benchtop diffractometer
 Absorption correction: multi-scan (SADABS; Sheldrick, 2008b)
 $T_{\min} = 0.955$, $T_{\max} = 0.985$

4796 measured reflections
 1413 independent reflections
 1226 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.041$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.037$
 $wR(F^2) = 0.099$
 $S = 1.05$
 1413 reflections
 185 parameters

1 restraint
 H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.15$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.14$ e Å⁻³

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{C8}-\text{H8B}\cdots\text{O3}^{\text{i}}$	0.97	2.62	3.551 (4)	161
$\text{C13}-\text{H13C}\cdots\text{O3}^{\text{ii}}$	0.96	2.63	3.533 (4)	156
$\text{C8}-\text{H8A}\cdots\text{O5}^{\text{iii}}$	0.97	2.44	3.309 (4)	149
$\text{C11}-\text{H11A}\cdots\text{O5}^{\text{iii}}$	0.96	2.70	3.662 (4)	178

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

Data collection: *GIS* (Bruker, 2010); cell refinement: *APEX2* (Bruker, 2010) and *SAINT* (Bruker, 2009); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008a); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008a); molecular graphics: *Mercury* (Macrae *et al.*, 2006); software used to prepare material for publication: *SHELXL97*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: IS2600).

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supplementary materials

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(1*S*,2*S*,6*S*,9*S*)-6-Methyl-5-oxobicyclo[4.4.0]decane-2,9-diyl diacetate

R. Aav, K. Lippur, M. Lopp and F. Werner

Comment

At 300 K the enantiopure compound (1*S*,2*S*,6*S*,9*S*)-6-Methyl-5-oxobicyclo[4.4.0]decane-2,9-diyl diacetate, (**I**), crystallizes in the chirodescriptive monoclinic space group *C*2 (No. 5) with one molecule in the asymmetric unit. Bond lengths and bond angles in the molecule are normal. The *trans*-fused cyclohexane rings both adopt chair conformation. The acetyl groups are inclined to the least-squares plane, defined by the carbon atoms of the cyclohexane rings, by $\sim 46.4^\circ$ (O2O3C12C13) and $\sim 51.2^\circ$ (O4O5C14C15), respectively (Fig. 1). The molecules are loosely held together in layers parallel to the *A*-plane with a repeating distance of $d_{100}/2 \sim 11.2 \text{ \AA}$, within which weak intra- and intermolecular hydrogen bonds (Steiner, 2002) occur (Fig. 2, Table 1). Between the layers only hydrophobic interactions are present.

Experimental

Enantiopure (**I**) was synthesized according to Aav *et al.* (2000). Single crystals were grown by slow evaporation of a solution of (**I**) in acetone/petrol ether.

Refinement

Owing to absence of significant anomalous scattering, Friedel pairs were merged and all f'' values were set to zero for the final refinement. The absolute structure was assigned from the synthetic procedure. Hydrogen atoms were included at calculated positions [$d(\text{C}-\text{H}) = 0.96$ (CH₃), 0.97 (CH₂) or 0.98 Å (CH)] and treated as riding on their base atoms, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ (CH₂ and CH) or $1.5U_{\text{eq}}(\text{C})$ (CH₃).

Figures

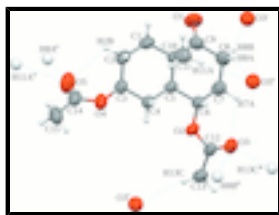


Fig. 1. Asymmetric unit in the crystal structure of (**I**). Displacement ellipsoids for non-H atoms are drawn at the 50% probability level. Cyan dashed lines indicate weak hydrogen bonds. [Symmetry codes: (i) $x, y, 1 + z$; (ii) $x, 1 + y, z$; (iii) $1/2 - x, 1/2 + y, -z$; (iv) $x, y, -1 + z$; (v) $1/2 - x, -1/2 + y, -z$; (vi) $x, -1 + y, z$.]

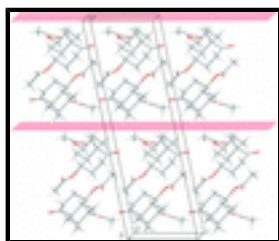


Fig. 2. Packing diagram of (**I**). Red planes indicate the boundaries of the layers within which weak hydrogen bonds occur. The unit cell is outlined.

(1S,2S,6S,9S)-\ 6-Methyl-5-oxobicyclo[4.4.0]decane-2,9-diyl diacetate

Crystal data

$C_{15}H_{22}O_5$	$F(000) = 608$
$M_r = 282.33$	$D_x = 1.238 \text{ Mg m}^{-3}$
Monoclinic, $C2$	Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$
Hall symbol: C 2y	Cell parameters from 2044 reflections
$a = 22.885 (5) \text{ \AA}$	$\theta = 2.4\text{--}23.9^\circ$
$b = 9.340 (2) \text{ \AA}$	$\mu = 0.09 \text{ mm}^{-1}$
$c = 7.2250 (13) \text{ \AA}$	$T = 300 \text{ K}$
$\beta = 101.280 (6)^\circ$	Needle, colorless
$V = 1514.5 (5) \text{ \AA}^3$	$0.50 \times 0.20 \times 0.16 \text{ mm}$
$Z = 4$	

Data collection

Bruker SMART X2S benchtop diffractometer	1413 independent reflections
Radiation source: XOS X-beam microfocus source doubly curved silicon crystal	1226 reflections with $I > 2\sigma(I)$
ω scans	$R_{\text{int}} = 0.041$
Absorption correction: multi-scan (SADABS; Sheldrick, 2008b)	$\theta_{\text{max}} = 25.0^\circ$, $\theta_{\text{min}} = 2.4^\circ$
$T_{\text{min}} = 0.955$, $T_{\text{max}} = 0.985$	$h = -27 \rightarrow 27$
4796 measured reflections	$k = -11 \rightarrow 11$
	$l = -7 \rightarrow 8$

Refinement

Refinement on F^2	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.037$	H-atom parameters constrained
$wR(F^2) = 0.099$	$w = 1/[\sigma^2(F_o^2) + (0.0528P)^2 + 0.0812P]$
$S = 1.05$	where $P = (F_o^2 + 2F_c^2)/3$
1413 reflections	$(\Delta/\sigma)_{\text{max}} < 0.001$
185 parameters	$\Delta\rho_{\text{max}} = 0.15 \text{ e \AA}^{-3}$
1 restraint	$\Delta\rho_{\text{min}} = -0.14 \text{ e \AA}^{-3}$
Primary atom site location: structure-invariant direct methods	Extinction correction: SHELXL97 (Sheldrick, 2008a), $F_c^* = kF_c[1 + 0.001x F_c^2 \lambda^3 / \sin(2\theta)]^{-1/4}$
	Extinction coefficient: 0.010 (2)

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)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.41782 (12)	0.1268 (3)	0.8403 (4)	0.0548 (7)
H1A	0.4505	0.1429	0.9460	0.066*
H1B	0.3854	0.0837	0.8891	0.066*
C2	0.43814 (13)	0.0228 (3)	0.7031 (5)	0.0603 (8)
H2A	0.4735	0.0603	0.6649	0.072*
H2B	0.4484	-0.0683	0.7656	0.072*
C3	0.38967 (12)	0.0004 (3)	0.5307 (4)	0.0507 (7)
H3	0.3559	-0.0484	0.5683	0.061*
C4	0.36824 (12)	0.1415 (3)	0.4352 (4)	0.0481 (6)
H4B	0.3359	0.1237	0.3292	0.058*
H4A	0.4005	0.1868	0.3878	0.058*
C5	0.34678 (11)	0.2411 (2)	0.5769 (4)	0.0417 (6)
H5	0.3154	0.1895	0.6241	0.050*
C6	0.31877 (12)	0.3794 (3)	0.4877 (4)	0.0460 (6)
H6	0.3485	0.4372	0.4405	0.055*
C7	0.29180 (14)	0.4638 (3)	0.6307 (4)	0.0619 (8)
H7B	0.2590	0.4099	0.6630	0.074*
H7A	0.2761	0.5536	0.5745	0.074*
C8	0.33767 (15)	0.4945 (3)	0.8099 (4)	0.0625 (8)
H8A	0.3653	0.5665	0.7822	0.075*
H8B	0.3174	0.5338	0.9044	0.075*
C9	0.37226 (13)	0.3646 (3)	0.8900 (4)	0.0559 (7)
O1	0.38133 (13)	0.3386 (3)	1.0571 (3)	0.0894 (8)
C10	0.39716 (11)	0.2710 (3)	0.7505 (4)	0.0458 (6)
C11	0.45030 (12)	0.3532 (4)	0.6993 (4)	0.0619 (7)
H11A	0.4377	0.4477	0.6568	0.093*
H11B	0.4645	0.3033	0.6006	0.093*
H11C	0.4817	0.3598	0.8085	0.093*
O2	0.27152 (7)	0.33569 (19)	0.3333 (2)	0.0510 (5)
C12	0.25230 (13)	0.4324 (3)	0.1969 (4)	0.0523 (7)
O3	0.26894 (10)	0.5536 (2)	0.2034 (3)	0.0678 (6)
C13	0.20704 (15)	0.3674 (4)	0.0446 (4)	0.0689 (9)
H13A	0.1906	0.4400	-0.0446	0.103*

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H13B	0.1758	0.3257	0.0982	0.103*
H13C	0.2254	0.2945	-0.0183	0.103*
O4	0.41045 (9)	-0.08472 (18)	0.3880 (3)	0.0610 (6)
C14	0.41311 (14)	-0.2269 (3)	0.4139 (5)	0.0673 (9)
O5	0.40094 (15)	-0.2845 (3)	0.5499 (5)	0.1017 (10)
C15	0.43333 (19)	-0.3023 (4)	0.2556 (6)	0.0896 (12)
H15A	0.4749	-0.3250	0.2920	0.134*
H15B	0.4272	-0.2414	0.1464	0.134*
H15C	0.4109	-0.3889	0.2263	0.134*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0477 (15)	0.0635 (17)	0.0480 (15)	-0.0007 (12)	-0.0035 (12)	0.0094 (13)
C2	0.0505 (16)	0.0594 (17)	0.0648 (19)	0.0099 (13)	-0.0041 (14)	0.0092 (14)
C3	0.0501 (15)	0.0431 (12)	0.0586 (17)	0.0034 (11)	0.0097 (13)	0.0039 (13)
C4	0.0499 (15)	0.0457 (13)	0.0460 (15)	0.0032 (11)	0.0029 (12)	0.0039 (11)
C5	0.0389 (12)	0.0430 (12)	0.0428 (14)	-0.0014 (10)	0.0071 (11)	0.0087 (11)
C6	0.0471 (13)	0.0471 (13)	0.0435 (14)	0.0033 (10)	0.0081 (12)	0.0023 (11)
C7	0.0649 (18)	0.0655 (18)	0.0561 (17)	0.0190 (14)	0.0137 (16)	0.0037 (15)
C8	0.080 (2)	0.0593 (16)	0.0500 (16)	0.0057 (15)	0.0167 (16)	-0.0031 (14)
C9	0.0597 (16)	0.0617 (17)	0.0464 (16)	-0.0076 (13)	0.0107 (13)	-0.0014 (13)
O1	0.128 (2)	0.0996 (19)	0.0414 (12)	0.0212 (18)	0.0181 (13)	0.0080 (13)
C10	0.0410 (13)	0.0527 (14)	0.0430 (14)	-0.0023 (11)	0.0067 (11)	0.0048 (11)
C11	0.0475 (14)	0.0716 (18)	0.0660 (18)	-0.0132 (14)	0.0093 (13)	-0.0024 (16)
O2	0.0517 (10)	0.0500 (10)	0.0470 (10)	0.0037 (8)	-0.0007 (8)	0.0103 (9)
C12	0.0601 (16)	0.0520 (15)	0.0464 (16)	0.0186 (13)	0.0140 (14)	0.0063 (13)
O3	0.0948 (17)	0.0498 (11)	0.0581 (13)	0.0096 (11)	0.0132 (11)	0.0094 (10)
C13	0.079 (2)	0.0689 (19)	0.0520 (17)	0.0173 (16)	-0.0041 (15)	0.0027 (15)
O4	0.0668 (13)	0.0433 (10)	0.0724 (15)	0.0070 (8)	0.0127 (11)	0.0004 (9)
C14	0.0613 (18)	0.0466 (16)	0.085 (3)	0.0011 (13)	-0.0078 (17)	0.0000 (16)
O5	0.138 (3)	0.0503 (12)	0.117 (2)	-0.0015 (14)	0.024 (2)	0.0167 (14)
C15	0.093 (3)	0.0586 (18)	0.106 (3)	0.0127 (18)	-0.007 (2)	-0.0224 (19)

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

C1—C2	1.524 (4)	C8—H8A	0.9700
C1—C10	1.528 (4)	C8—H8B	0.9700
C1—H1A	0.9700	C9—O1	1.209 (3)
C1—H1B	0.9700	C9—C10	1.526 (4)
C2—C3	1.512 (4)	C10—C11	1.543 (4)
C2—H2A	0.9700	C11—H11A	0.9600
C2—H2B	0.9700	C11—H11B	0.9600
C3—O4	1.453 (3)	C11—H11C	0.9600
C3—C4	1.523 (4)	O2—C12	1.346 (3)
C3—H3	0.9800	C12—O3	1.193 (4)
C4—C5	1.533 (3)	C12—O3	1.193 (4)
C4—H4B	0.9700	C12—C13	1.485 (4)
C4—H4A	0.9700	C13—H13A	0.9600

C5—C6	1.527 (3)	C13—H13B	0.9600
C5—C10	1.554 (3)	C13—H13C	0.9600
C5—H5	0.9800	O4—C14	1.340 (4)
C6—O2	1.452 (3)	C14—O5	1.200 (4)
C6—C7	1.523 (4)	C14—O5	1.200 (4)
C6—H6	0.9800	C14—C15	1.492 (5)
C7—C8	1.526 (4)	C15—H15A	0.9600
C7—H7B	0.9700	C15—H15B	0.9600
C7—H7A	0.9700	C15—H15C	0.9600
C8—C9	1.502 (4)		
C2—C1—C10	113.2 (2)	C7—C8—H8A	108.9
C2—C1—H1A	108.9	C9—C8—H8B	108.9
C10—C1—H1A	108.9	C7—C8—H8B	108.9
C2—C1—H1B	108.9	H8A—C8—H8B	107.7
C10—C1—H1B	108.9	O1—C9—C8	121.5 (3)
H1A—C1—H1B	107.8	O1—C9—C10	122.1 (3)
C3—C2—C1	110.9 (2)	C8—C9—C10	116.4 (2)
C3—C2—H2A	109.5	C9—C10—C1	110.4 (2)
C1—C2—H2A	109.5	C9—C10—C11	106.7 (2)
C3—C2—H2B	109.5	C1—C10—C11	110.3 (2)
C1—C2—H2B	109.5	C9—C10—C5	108.8 (2)
H2A—C2—H2B	108.1	C1—C10—C5	107.7 (2)
O4—C3—C2	111.8 (2)	C11—C10—C5	113.0 (2)
O4—C3—C4	105.9 (2)	C10—C11—H11A	109.5
C2—C3—C4	111.9 (2)	C10—C11—H11B	109.5
O4—C3—H3	109.1	H11A—C11—H11B	109.5
C2—C3—H3	109.1	C10—C11—H11C	109.5
C4—C3—H3	109.1	H11A—C11—H11C	109.5
C3—C4—C5	109.8 (2)	H11B—C11—H11C	109.5
C3—C4—H4B	109.7	C12—O2—C6	117.5 (2)
C5—C4—H4B	109.7	O3—C12—O2	123.5 (3)
C3—C4—H4A	109.7	O3—C12—O2	123.5 (3)
C5—C4—H4A	109.7	O3—C12—C13	126.0 (3)
H4B—C4—H4A	108.2	O3—C12—C13	126.0 (3)
C6—C5—C4	113.18 (19)	O2—C12—C13	110.4 (3)
C6—C5—C10	111.9 (2)	C12—C13—H13A	109.5
C4—C5—C10	111.34 (19)	C12—C13—H13B	109.5
C6—C5—H5	106.7	H13A—C13—H13B	109.5
C4—C5—H5	106.7	C12—C13—H13C	109.5
C10—C5—H5	106.7	H13A—C13—H13C	109.5
O2—C6—C7	109.1 (2)	H13B—C13—H13C	109.5
O2—C6—C5	105.94 (19)	C14—O4—C3	117.1 (3)
C7—C6—C5	110.1 (2)	O5—C14—O4	123.2 (3)
O2—C6—H6	110.5	O5—C14—O4	123.2 (3)
C7—C6—H6	110.5	O5—C14—C15	124.9 (3)
C5—C6—H6	110.5	O5—C14—C15	124.9 (3)
C6—C7—C8	111.7 (2)	O4—C14—C15	111.9 (3)
C6—C7—H7B	109.3	C14—C15—H15A	109.5
C8—C7—H7B	109.3	C14—C15—H15B	109.5

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C6—C7—H7A	109.3	H15A—C15—H15B	109.5
C8—C7—H7A	109.3	C14—C15—H15C	109.5
H7B—C7—H7A	107.9	H15A—C15—H15C	109.5
C9—C8—C7	113.5 (3)	H15B—C15—H15C	109.5
C9—C8—H8A	108.9		

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

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
C7—H7A \cdots O3	0.97	2.65	3.143 (4)	112
C8—H8B \cdots O3 ⁱ	0.97	2.62	3.551 (4)	161
C13—H13C \cdots O3 ⁱⁱ	0.96	2.63	3.533 (4)	156
C2—H2B \cdots O5	0.97	2.65	3.134 (4)	111
C8—H8A \cdots O5 ⁱⁱⁱ	0.97	2.44	3.309 (4)	149
C11—H11A \cdots O5 ⁱⁱⁱ	0.96	2.70	3.662 (4)	178

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

Fig. 1

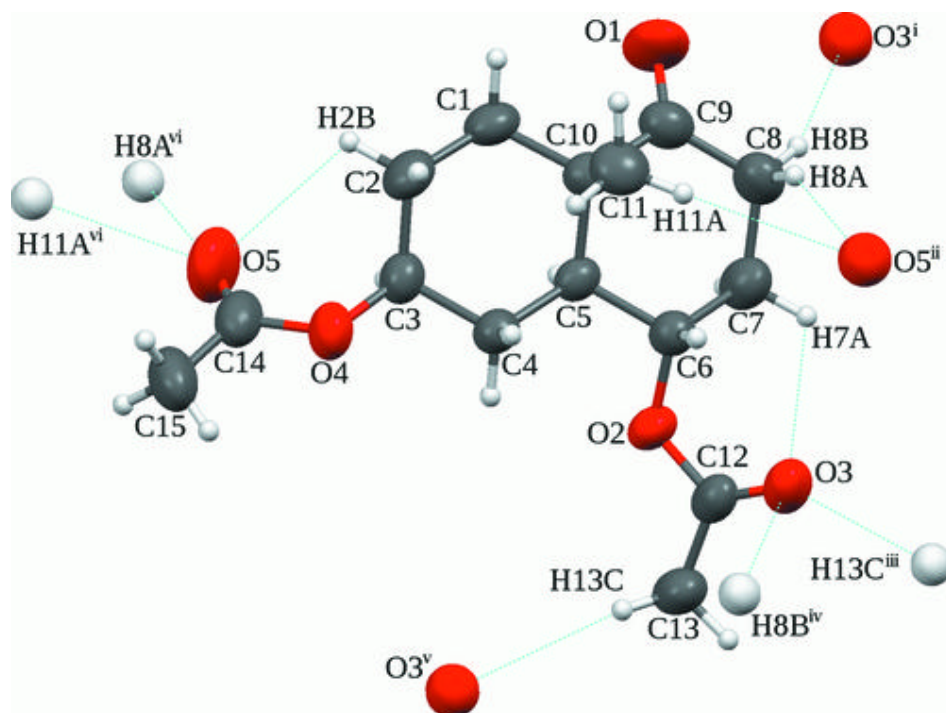


Fig. 2

