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(2*R*,3*S*)-2-Benzyl-3-(2,3,4,6-tetra-*O*-acetyl-β-D-glucopyranosyloxy)butanolide

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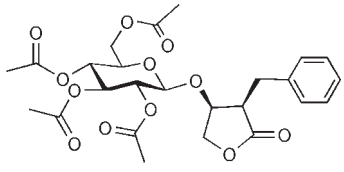
Key indicators: single-crystal X-ray study; T = 153 K; mean σ(C–C) = 0.003 Å; R factor = 0.036; wR factor = 0.102; data-to-parameter ratio = 12.3.

The title compound, $C_{25}H_{30}O_{12}$, which demonstrates a significant hepatoprotective effect, has comparable geometrical parameters to those of similar compounds. The absolute configuration of the title compound, *viz*. 2*R*,3*S*, was identified from the Flack parameter of 0.05 (17) and the Hooft parameter of 0.04 (6).

Related literature

For the hepatoprotective effect of the title compound, see: Du & Irinon (2008). For bond-length data, see: Allen *et al.* (1987). For the Hooft parameter, see: Hooft *et al.*(2008). For details of the preparation, see: Saito *et al.* (1992); Kazumasa *et al.* (2000); Schmidt (1986); Corey & Venkateswarlu (1972); Fernandez *et al.* (1997).



Experimental

Crystal data

$C_{25}H_{30}O_{12}$
M_r = 522.49
Orthorhombic, $P2_12_12_1$
a = 10.6142 (2) Å
b = 11.0984 (2) Å
c = 22.9714 (3) Å
V = 2706.05 (8) Å^3
Z = 4
Cu $K\alpha$ radiation
μ = 0.87 mm^{-1}
T = 153 K
0.15 × 0.15 × 0.10 mm

Data collection

Mac dip 2030b diffractometer
Absorption correction: multi-scan (*SADABS*; Sheldrick, 2003)
T_{min} = 0.880, T_{max} = 0.918
7246 measured reflections
4102 independent reflections
4015 reflections with $I > 2\sigma(I)$
R_{int} = 0.016

Refinement

$R[F^2 > 2\sigma(F^2)]$ = 0.036
$wR(F^2)$ = 0.102
S = 1.07
4102 reflections
334 parameters
H-atom parameters constrained
$\Delta\rho_{max}$ = 0.34 e Å^{-3}
$\Delta\rho_{min}$ = −0.22 e Å^{-3}
Absolute structure: Flack (1983), 1417 Friedel pairs
Flack parameter: 0.05 (17)

Data collection: *DENZO* (Otwinowski & Minor, 1997); cell refinement: *SCALEPACK* (Otwinowski & Minor, 1997); data reduction: *SCALEPACK*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEPII* (Johnson, 1976) and *PLATON* (Spek, 2009); 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: VM2019).

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supplementary materials

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(2*R*,3*S*)-2-Benzyl-3-(2,3,4,6-tetra-*O*-acetyl-β-D-glucopyranosyloxy)butanolide

F. Zhang, J. Zhong, B. Han, D. Yin and H. Huang

Comment

The title compound was found to have a significant hepatoprotective effect, comparable to the natural product Goodyeroside A (Du & Irinon, 2008). Recently, the compound was successfully crystallized from a petroleum ether/EtOAc mixture, yielding crystals suitable for X-ray analysis.

Fig.1 shows the molecular structure of the title compound with atomic numbering scheme. The bond lengths and angles are in agreement with reported literature values (Table 1) (Allen *et al.*, 1987). The benzene ring (C6—C11) is essentially planar, with r.m.s deviations of 0.0078 (14) Å. The five-membered ring (C1—C4/O2) has an envelope conformation with the C3 atom out of plane and the six-membered ring (C1'-C5'/O4) is in its chair conformation. The dihedral angles between the various rings in the title compound are as follows, where the first atom is used to identify its five- or six- membered ring: C1/C6 112.9 (1)°; C1/C1' 110.1 (1)°; C6/C1' 13.1 (1)°.

In order to determine the absolute configuration of the title compound, the data collection was performed using Cu Kα. The absolute configuration is confirmed by the Flack parameter 0.05 (17) and Hooft parameter 0.04 (6) (Flack, 1983; Hooft *et al.*, 2008).

Experimental

To a solution of Ethyl (2*R*,3*S*)-3-(2,3,4,6-tetra-*O*-acetyl-β-D-glucopyranosyloxy)-2- benzyl-4-hydroxybutanoate (1.0 equiv) in 1,4-dioxane (10 ml) was added 4-methylbenzenesulfonic acid (1.0 equiv), then the solution was stirred at room temperature for 2 days. After removal of the solvent in vacuo, the residue was purified by column chromatography. White crystals suitable for X-ray analysis were obtained by slow evaportion of a petroleum ether/EtOAc solution over a period of two weeks (Saito *et al.*, 1992; Kazumasa *et al.*, 2000; Schmidt, 1986; Corey & Venkateswarlu, 1972; Fernandez *et al.*, 1997).

Refinement

All H atoms were placed in geometrically idealised positions and constrained to ride on their parent atoms with C—H distances in the range of 0.95-1.00 Å, with a displacement parameter U_{iso} set to 1.2 (CH and CH_2) or 1.5(CH_3) times U_{eq} of the parent atom.

Figures

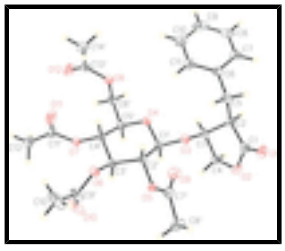


Fig. 1. View of the title compound showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 30% probability level.

(2*R*,3*S*)-2-Benzyl-3-(2,3,4,6-tetra-*O*-acetyl-β-D- glucopyranosyloxy)butanolide

Crystal data

$C_{25}H_{30}O_{12}$	$F(000) = 1104$
$M_r = 522.49$	$D_x = 1.282$ Mg m^{-3}
Orthorhombic, $P2_12_12_1$	Cu $K\alpha$ radiation, $\lambda = 1.54178$ Å
Hall symbol: P 2ac 2ab	Cell parameters from 5304 reflections
$a = 10.6142$ (2) Å	$\theta = 4.0$–$65.7°$
$b = 11.0984$ (2) Å	$\mu = 0.87$ mm^{-1}
$c = 22.9714$ (3) Å	$T = 153$ K
$V = 2706.05$ (8) Å^3	Block, white
$Z = 4$	0.15 × 0.15 × 0.10 mm

Data collection

Mac dip 2030b diffractometer	4102 independent reflections
Radiation source: rotating anode	4015 reflections with $I > 2\sigma(I)$
graphite	$R_{int} = 0.016$
Detector resolution: 0 pixels mm^{-1}	$\theta_{max} = 66.0°$, $\theta_{min} = 4.4°$
ω scans	$h = -12 \rightarrow 9$
Absorption correction: multi-scan (SADABS; Sheldrick, 2003)	$k = -12 \rightarrow 12$
$T_{min} = 0.880$, $T_{max} = 0.918$	$l = -26 \rightarrow 26$
7246 measured reflections	

Refinement

Refinement on F^2	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.036$	H-atom parameters constrained
$wR(F^2) = 0.102$	$w = 1/[\sigma^2(F_o^2) + (0.0654P)^2 + 0.5803P]$ where $P = (F_o^2 + 2F_c^2)/3$
$S = 1.07$	$(\Delta/\sigma)_{max} < 0.001$

4102 reflections	$\Delta\rho_{max}$ = 0.34 e Å^{-3}
334 parameters	$\Delta\rho_{min}$ = −0.22 e Å^{-3}
0 restraints	Absolute structure: Flack (1983), 1417 Friedel pairs
Primary atom site location: structure-invariant direct methods	Flack parameter: 0.05 (17)

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 $F^2 > \sigma(F^2)$ 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 (Å^2)

	x	*y*	*z*	U_{iso}*/U_{eq}
O1	0.47650 (16)	0.31851 (14)	0.32119 (7)	0.0442 (4)
O2	0.58032 (14)	0.48590 (14)	0.29725 (6)	0.0403 (3)
O3	0.49559 (12)	0.47669 (12)	0.17568 (6)	0.0311 (3)
O4	0.44988 (12)	0.58270 (12)	0.09414 (6)	0.0300 (3)
O5	0.73101 (12)	0.40529 (12)	0.12561 (6)	0.0319 (3)
O6	0.71425 (13)	0.39829 (13)	0.00188 (6)	0.0351 (3)
O7	0.61363 (13)	0.62268 (12)	−0.04400 (6)	0.0325 (3)
O8	0.29897 (12)	0.61966 (13)	−0.00230 (6)	0.0353 (3)
O9	0.64011 (15)	0.22826 (13)	0.14807 (7)	0.0427 (4)
O10	0.89389 (19)	0.49770 (18)	−0.01722 (12)	0.0784 (7)
O11	0.5236 (2)	0.4937 (2)	−0.10623 (7)	0.0675 (6)
O12	0.2865 (2)	0.7043 (2)	−0.09013 (8)	0.0739 (7)
C1	0.4778 (2)	0.41371 (19)	0.29662 (9)	0.0341 (4)
C2	0.37455 (19)	0.47400 (18)	0.26180 (8)	0.0306 (4)
H2A	0.3281	0.5283	0.2891	0.037*
C3	0.45018 (18)	0.55428 (17)	0.22133 (8)	0.0297 (4)
H3A	0.4010	0.6247	0.2064	0.036*
C4	0.5581 (2)	0.59050 (19)	0.26080 (9)	0.0351 (4)
H4A	0.6341	0.6103	0.2377	0.042*
H4B	0.5350	0.6613	0.2847	0.042*
C5	0.2771 (2)	0.38840 (18)	0.23518 (9)	0.0346 (4)
H5A	0.3174	0.3409	0.2038	0.042*
H5B	0.2477	0.3314	0.2654	0.042*
C6	0.16563 (19)	0.45524 (18)	0.21066 (10)	0.0354 (4)
C7	0.0715 (2)	0.4974 (2)	0.24720 (10)	0.0421 (5)
H7A	0.0756	0.4808	0.2877	0.050*
C8	−0.0285 (2)	0.5636 (2)	0.22519 (15)	0.0569 (7)

H8A	−0.0916	0.5928	0.2509	0.068*
C9	−0.0377 (3)	0.5875 (2)	0.16642 (15)	0.0616 (8)
H9A	−0.1064	0.6328	0.1515	0.074*
C10	0.0552 (3)	0.5443 (3)	0.12938 (13)	0.0606 (8)
H10A	0.0493	0.5591	0.0887	0.073*
C11	0.1569 (2)	0.4795 (2)	0.15118 (10)	0.0456 (5)
H11A	0.2207	0.4516	0.1255	0.055*
C1'	0.54911 (18)	0.53583 (18)	0.12866 (8)	0.0294 (4)
H1'A	0.6062	0.6018	0.1421	0.035*
C2'	0.62123 (18)	0.44333 (18)	0.09365 (8)	0.0283 (4)
H2'A	0.5661	0.3723	0.0854	0.034*
C3'	0.66843 (17)	0.49655 (17)	0.03700 (8)	0.0281 (4)
H3'A	0.7380	0.5551	0.0448	0.034*
C4'	0.56170 (18)	0.55880 (17)	0.00501 (8)	0.0288 (4)
H4'A	0.4997	0.4974	−0.0089	0.035*
C5'	0.49602 (18)	0.64833 (18)	0.04534 (8)	0.0299 (4)
H5'A	0.5582	0.7100	0.0589	0.036*
C6'	0.38565 (19)	0.71066 (19)	0.01696 (9)	0.0359 (5)
H6'A	0.3439	0.7650	0.0452	0.043*
H6'B	0.4148	0.7593	−0.0166	0.043*
C7'	0.7291 (2)	0.29459 (19)	0.15016 (9)	0.0351 (4)
C8'	0.8527 (3)	0.2666 (2)	0.17810 (13)	0.0547 (6)
H8'A	0.8493	0.1860	0.1955	0.082*
H8'B	0.8703	0.3262	0.2085	0.082*
H8'C	0.9195	0.2692	0.1487	0.082*
C9'	0.8281 (2)	0.4122 (2)	−0.02386 (12)	0.0516 (6)
C10'	0.8550 (4)	0.3043 (3)	−0.06145 (19)	0.0899 (12)
H10B	0.9370	0.3143	−0.0804	0.135*
H10C	0.7892	0.2967	−0.0912	0.135*
H10D	0.8562	0.2315	−0.0373	0.135*
C11'	0.5904 (2)	0.5778 (2)	−0.09765 (9)	0.0385 (5)
C12'	0.6610 (2)	0.6479 (2)	−0.14304 (10)	0.0464 (5)
H12A	0.6431	0.6142	−0.1816	0.070*
H12B	0.7516	0.6428	−0.1353	0.070*
H12C	0.6343	0.7324	−0.1419	0.070*
C13'	0.26186 (19)	0.6221 (2)	−0.05781 (9)	0.0403 (5)
C14'	0.1842 (2)	0.5152 (3)	−0.07186 (11)	0.0526 (6)
H14A	0.1585	0.5186	−0.1128	0.079*
H14B	0.1091	0.5144	−0.0470	0.079*
H14C	0.2334	0.4419	−0.0650	0.079*

Atomic displacement parameters (Å^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0555 (9)	0.0361 (8)	0.0410 (8)	0.0107 (7)	−0.0020 (7)	0.0053 (7)
O2	0.0368 (8)	0.0424 (9)	0.0418 (7)	0.0029 (7)	−0.0093 (6)	−0.0006 (7)
O3	0.0359 (7)	0.0277 (7)	0.0297 (6)	−0.0020 (5)	0.0038 (5)	−0.0030 (6)
O4	0.0273 (7)	0.0310 (7)	0.0318 (6)	0.0005 (6)	0.0018 (5)	−0.0016 (6)

O5	0.0276 (7)	0.0281 (7)	0.0401 (7)	−0.0025 (6)	−0.0058 (6)	0.0042 (6)
O6	0.0340 (7)	0.0320 (7)	0.0393 (7)	0.0047 (6)	0.0059 (6)	−0.0049 (6)
O7	0.0321 (7)	0.0342 (7)	0.0313 (6)	−0.0013 (6)	0.0026 (6)	0.0017 (6)
O8	0.0294 (7)	0.0396 (8)	0.0369 (7)	−0.0006 (6)	−0.0016 (6)	0.0059 (6)
O9	0.0455 (9)	0.0308 (8)	0.0518 (8)	−0.0081 (7)	−0.0055 (7)	0.0073 (7)
O10	0.0542 (11)	0.0501 (12)	0.1310 (19)	−0.0041 (10)	0.0518 (12)	−0.0067 (12)
O11	0.0873 (14)	0.0811 (15)	0.0342 (8)	−0.0362 (13)	−0.0059 (8)	−0.0060 (9)
O12	0.0730 (13)	0.0984 (17)	0.0504 (10)	−0.0315 (12)	−0.0147 (9)	0.0310 (11)
C1	0.0384 (11)	0.0333 (11)	0.0305 (9)	0.0050 (9)	0.0005 (8)	−0.0050 (9)
C2	0.0327 (10)	0.0287 (10)	0.0303 (9)	0.0043 (8)	0.0020 (8)	−0.0028 (8)
C3	0.0300 (10)	0.0271 (10)	0.0320 (9)	0.0022 (8)	−0.0005 (8)	−0.0042 (8)
C4	0.0331 (10)	0.0339 (11)	0.0383 (10)	−0.0011 (8)	−0.0029 (8)	−0.0039 (9)
C5	0.0345 (10)	0.0280 (10)	0.0413 (10)	−0.0006 (9)	0.0030 (9)	−0.0007 (9)
C6	0.0317 (10)	0.0280 (10)	0.0464 (11)	−0.0061 (8)	−0.0026 (9)	−0.0024 (9)
C7	0.0312 (10)	0.0416 (12)	0.0534 (12)	−0.0050 (9)	−0.0025 (9)	−0.0091 (11)
C8	0.0311 (12)	0.0459 (14)	0.094 (2)	−0.0005 (10)	−0.0113 (12)	−0.0126 (14)
C9	0.0429 (14)	0.0391 (13)	0.103 (2)	−0.0070 (11)	−0.0301 (15)	0.0106 (15)
C10	0.0598 (17)	0.0572 (16)	0.0650 (16)	−0.0182 (14)	−0.0268 (14)	0.0211 (14)
C11	0.0473 (12)	0.0424 (13)	0.0471 (12)	−0.0111 (11)	−0.0049 (10)	0.0031 (10)
C1'	0.0280 (9)	0.0284 (10)	0.0317 (9)	−0.0053 (8)	−0.0001 (8)	−0.0017 (8)
C2'	0.0234 (9)	0.0271 (10)	0.0342 (9)	−0.0028 (7)	−0.0031 (8)	0.0009 (8)
C3'	0.0251 (9)	0.0255 (9)	0.0337 (9)	−0.0003 (7)	0.0034 (7)	−0.0034 (8)
C4'	0.0273 (9)	0.0280 (10)	0.0310 (9)	−0.0028 (7)	0.0018 (8)	−0.0006 (8)
C5'	0.0277 (10)	0.0282 (10)	0.0338 (9)	−0.0011 (7)	0.0027 (8)	−0.0001 (8)
C6'	0.0309 (10)	0.0329 (11)	0.0439 (11)	0.0045 (9)	0.0021 (9)	0.0018 (9)
C7'	0.0404 (11)	0.0284 (10)	0.0366 (10)	0.0008 (9)	−0.0039 (9)	0.0014 (8)
C8'	0.0558 (15)	0.0383 (13)	0.0701 (15)	−0.0020 (11)	−0.0260 (13)	0.0120 (12)
C9'	0.0476 (13)	0.0386 (13)	0.0686 (15)	0.0099 (11)	0.0254 (12)	0.0046 (12)
C10'	0.087 (2)	0.0606 (18)	0.122 (3)	0.0138 (17)	0.060 (2)	−0.014 (2)
C11'	0.0371 (12)	0.0449 (13)	0.0335 (10)	0.0044 (10)	−0.0032 (9)	−0.0009 (9)
C12'	0.0471 (13)	0.0555 (14)	0.0365 (10)	0.0082 (11)	0.0073 (10)	0.0059 (10)
C13'	0.0272 (10)	0.0571 (14)	0.0367 (10)	0.0039 (9)	−0.0007 (9)	0.0101 (10)
C14'	0.0434 (13)	0.0680 (17)	0.0464 (12)	−0.0034 (12)	−0.0112 (10)	0.0012 (12)

Geometric parameters (Å, °)

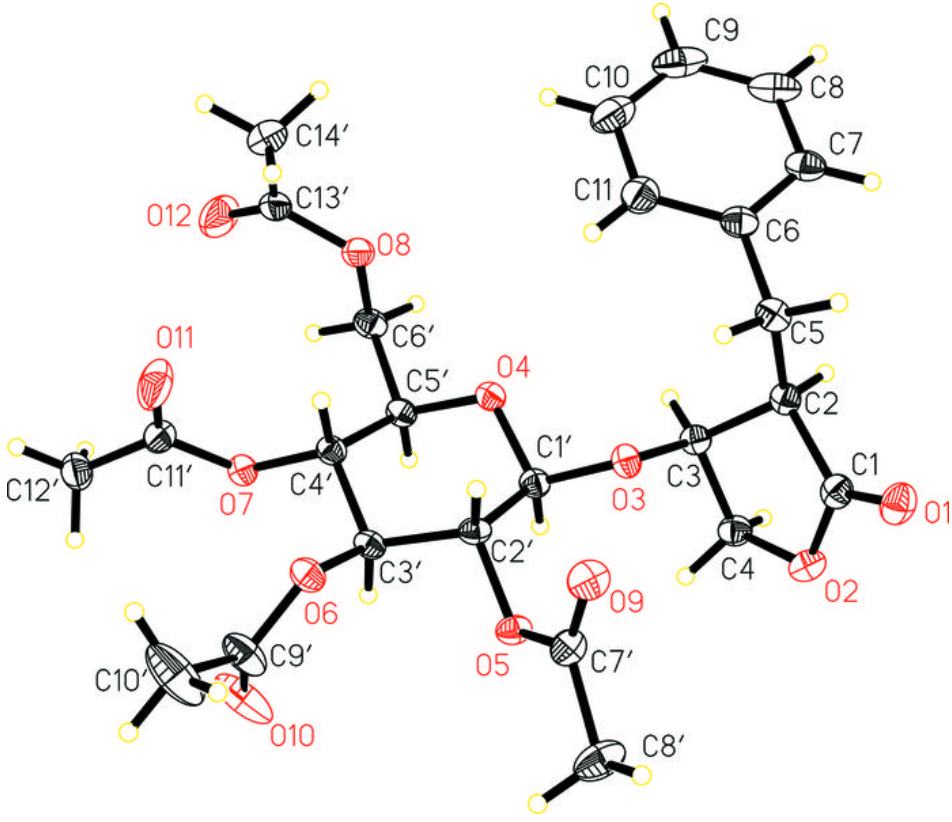
O1—C1	1.198 (3)	C8—H8A	0.9500
O2—C1	1.351 (3)	C9—C10	1.388 (5)
O2—C4	1.451 (3)	C9—H9A	0.9500
O3—C1'	1.386 (2)	C10—C11	1.390 (4)
O3—C3	1.440 (2)	C10—H10A	0.9500
O4—C1'	1.417 (2)	C11—H11A	0.9500
O4—C5'	1.424 (2)	C1'—C2'	1.512 (3)
O5—C7'	1.352 (3)	C1'—H1'A	1.0000
O5—C2'	1.440 (2)	C2'—C3'	1.514 (3)
O6—C9'	1.355 (3)	C2'—H2'A	1.0000
O6—C3'	1.441 (2)	C3'—C4'	1.517 (3)
O7—C11'	1.352 (2)	C3'—H3'A	1.0000
O7—C4'	1.440 (2)	C4'—C5'	1.527 (3)

O8—C13'	1.335 (3)	C4'—H4'A	1.0000
O8—C6'	1.436 (3)	C5'—C6'	1.509 (3)
O9—C7'	1.199 (3)	C5'—H5'A	1.0000
O10—C9'	1.188 (3)	C6'—H6'A	0.9900
O11—C11'	1.189 (3)	C6'—H6'B	0.9900
O12—C13'	1.205 (3)	C7'—C8'	1.493 (3)
C1—C2	1.513 (3)	C8'—H8'A	0.9800
C2—C3	1.517 (3)	C8'—H8'B	0.9800
C2—C5	1.532 (3)	C8'—H8'C	0.9800
C2—H2A	1.0000	C9'—C10'	1.504 (4)
C3—C4	1.515 (3)	C10'—H10B	0.9800
C3—H3A	1.0000	C10'—H10C	0.9800
C4—H4A	0.9900	C10'—H10D	0.9800
C4—H4B	0.9900	C11'—C12'	1.501 (3)
C5—C6	1.506 (3)	C12'—H12A	0.9800
C5—H5A	0.9900	C12'—H12B	0.9800
C5—H5B	0.9900	C12'—H12C	0.9800
C6—C7	1.386 (3)	C13'—C14'	1.480 (4)
C6—C11	1.396 (3)	C14'—H14A	0.9800
C7—C8	1.387 (3)	C14'—H14B	0.9800
C7—H7A	0.9500	C14'—H14C	0.9800
C8—C9	1.379 (5)		
C1—O2—C4	109.73 (15)	C1'—C2'—H2'A	109.8
C1'—O3—C3	114.93 (15)	C3'—C2'—H2'A	109.8
C1'—O4—C5'	111.88 (14)	O6—C3'—C2'	107.32 (15)
C7'—O5—C2'	117.84 (15)	O6—C3'—C4'	108.99 (15)
C9'—O6—C3'	117.33 (17)	C2'—C3'—C4'	110.30 (15)
C11'—O7—C4'	117.49 (16)	O6—C3'—H3'A	110.1
C13'—O8—C6'	117.98 (17)	C2'—C3'—H3'A	110.1
O1—C1—O2	121.81 (19)	C4'—C3'—H3'A	110.1
O1—C1—C2	129.1 (2)	O7—C4'—C3'	108.49 (15)
O2—C1—C2	109.08 (17)	O7—C4'—C5'	109.19 (15)
C1—C2—C3	101.56 (16)	C3'—C4'—C5'	110.08 (15)
C1—C2—C5	115.21 (17)	O7—C4'—H4'A	109.7
C3—C2—C5	118.49 (16)	C3'—C4'—H4'A	109.7
C1—C2—H2A	106.9	C5'—C4'—H4'A	109.7
C3—C2—H2A	106.9	O4—C5'—C6'	107.93 (16)
C5—C2—H2A	106.9	O4—C5'—C4'	107.57 (15)
O3—C3—C4	109.95 (16)	C6'—C5'—C4'	112.99 (16)
O3—C3—C2	105.78 (15)	O4—C5'—H5'A	109.4
C4—C3—C2	100.91 (16)	C6'—C5'—H5'A	109.4
O3—C3—H3A	113.1	C4'—C5'—H5'A	109.4
C4—C3—H3A	113.1	O8—C6'—C5'	107.95 (16)
C2—C3—H3A	113.1	O8—C6'—H6'A	110.1
O2—C4—C3	104.82 (16)	C5'—C6'—H6'A	110.1
O2—C4—H4A	110.8	O8—C6'—H6'B	110.1
C3—C4—H4A	110.8	C5'—C6'—H6'B	110.1
O2—C4—H4B	110.8	H6'A—C6'—H6'B	108.4
C3—C4—H4B	110.8	O9—C7'—O5	123.58 (19)

H4A—C4—H4B	108.9	O9—C7'—C8'	125.6 (2)
C6—C5—C2	111.99 (16)	O5—C7'—C8'	110.83 (18)
C6—C5—H5A	109.2	C7'—C8'—H8'A	109.5
C2—C5—H5A	109.2	C7'—C8'—H8'B	109.5
C6—C5—H5B	109.2	H8'A—C8'—H8'B	109.5
C2—C5—H5B	109.2	C7'—C8'—H8'C	109.5
H5A—C5—H5B	107.9	H8'A—C8'—H8'C	109.5
C7—C6—C11	118.7 (2)	H8'B—C8'—H8'C	109.5
C7—C6—C5	120.4 (2)	O10—C9'—O6	124.0 (2)
C11—C6—C5	120.9 (2)	O10—C9'—C10'	126.8 (2)
C8—C7—C6	120.6 (2)	O6—C9'—C10'	109.2 (2)
C8—C7—H7A	119.7	C9'—C10'—H10B	109.5
C6—C7—H7A	119.7	C9'—C10'—H10C	109.5
C9—C8—C7	120.9 (3)	H10B—C10'—H10C	109.5
C9—C8—H8A	119.6	C9'—C10'—H10D	109.5
C7—C8—H8A	119.6	H10B—C10'—H10D	109.5
C8—C9—C10	118.9 (2)	H10C—C10'—H10D	109.5
C8—C9—H9A	120.5	O11—C11'—O7	123.3 (2)
C10—C9—H9A	120.5	O11—C11'—C12'	126.1 (2)
C9—C10—C11	120.6 (3)	O7—C11'—C12'	110.57 (19)
C9—C10—H10A	119.7	C11'—C12'—H12A	109.5
C11—C10—H10A	119.7	C11'—C12'—H12B	109.5
C10—C11—C6	120.3 (3)	H12A—C12'—H12B	109.5
C10—C11—H11A	119.9	C11'—C12'—H12C	109.5
C6—C11—H11A	119.9	H12A—C12'—H12C	109.5
O3—C1'—O4	107.78 (15)	H12B—C12'—H12C	109.5
O3—C1'—C2'	107.49 (16)	O12—C13'—O8	122.7 (2)
O4—C1'—C2'	109.14 (14)	O12—C13'—C14'	126.4 (2)
O3—C1'—H1'A	110.8	O8—C13'—C14'	110.89 (19)
O4—C1'—H1'A	110.8	C13'—C14'—H14A	109.5
C2'—C1'—H1'A	110.8	C13'—C14'—H14B	109.5
O5—C2'—C1'	109.72 (15)	H14A—C14'—H14B	109.5
O5—C2'—C3'	106.54 (15)	C13'—C14'—H14C	109.5
C1'—C2'—C3'	111.08 (16)	H14A—C14'—H14C	109.5
O5—C2'—H2'A	109.8	H14B—C14'—H14C	109.5
C4—O2—C1—O1	176.02 (19)	O3—C1'—C2'—O5	−70.08 (18)
C4—O2—C1—C2	−4.4 (2)	O4—C1'—C2'—O5	173.29 (14)
O1—C1—C2—C3	−154.5 (2)	O3—C1'—C2'—C3'	172.38 (15)
O2—C1—C2—C3	25.89 (19)	O4—C1'—C2'—C3'	55.8 (2)
O1—C1—C2—C5	−25.2 (3)	C9'—O6—C3'—C2'	−132.97 (19)
O2—C1—C2—C5	155.26 (16)	C9'—O6—C3'—C4'	107.6 (2)
C1'—O3—C3—C4	−82.37 (19)	O5—C2'—C3'—O6	71.25 (18)
C1'—O3—C3—C2	169.45 (14)	C1'—C2'—C3'—O6	−169.29 (14)
C1—C2—C3—O3	79.21 (17)	O5—C2'—C3'—C4'	−170.14 (15)
C5—C2—C3—O3	−48.1 (2)	C1'—C2'—C3'—C4'	−50.7 (2)
C1—C2—C3—C4	−35.34 (18)	C11'—O7—C4'—C3'	108.25 (18)
C5—C2—C3—C4	−162.61 (17)	C11'—O7—C4'—C5'	−131.76 (18)
C1—O2—C4—C3	−19.3 (2)	O6—C3'—C4'—O7	−70.62 (18)
O3—C3—C4—O2	−77.30 (19)	C2'—C3'—C4'—O7	171.80 (15)

C2—C3—C4—O2	34.09 (19)	O6—C3'—C4'—C5'	169.95 (14)
C1—C2—C5—C6	169.70 (17)	C2'—C3'—C4'—C5'	52.4 (2)
C3—C2—C5—C6	−69.8 (2)	C1'—O4—C5'—C6'	−171.17 (16)
C2—C5—C6—C7	−78.9 (2)	C1'—O4—C5'—C4'	66.61 (18)
C2—C5—C6—C11	98.9 (2)	O7—C4'—C5'—O4	−177.99 (14)
C11—C6—C7—C8	−0.8 (3)	C3'—C4'—C5'—O4	−58.99 (19)
C5—C6—C7—C8	177.1 (2)	O7—C4'—C5'—C6'	63.0 (2)
C6—C7—C8—C9	0.9 (4)	C3'—C4'—C5'—C6'	−178.01 (16)
C7—C8—C9—C10	−0.1 (4)	C13'—O8—C6'—C5'	−127.09 (18)
C8—C9—C10—C11	−1.0 (4)	O4—C5'—C6'—O8	−61.5 (2)
C9—C10—C11—C6	1.1 (4)	C4'—C5'—C6'—O8	57.4 (2)
C7—C6—C11—C10	−0.3 (3)	C2'—O5—C7'—O9	−2.1 (3)
C5—C6—C11—C10	−178.1 (2)	C2'—O5—C7'—C8'	176.30 (19)
C3—O3—C1'—O4	−77.32 (18)	C3'—O6—C9'—O10	3.2 (4)
C3—O3—C1'—C2'	165.16 (15)	C3'—O6—C9'—C10'	−176.3 (2)
C5'—O4—C1'—O3	178.36 (14)	C4'—O7—C11'—O11	4.8 (3)
C5'—O4—C1'—C2'	−65.20 (19)	C4'—O7—C11'—C12'	−174.66 (18)
C7'—O5—C2'—C1'	106.05 (19)	C6'—O8—C13'—O12	−8.6 (3)
C7'—O5—C2'—C3'	−133.61 (17)	C6'—O8—C13'—C14'	173.88 (18)

Fig. 1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