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3β,11α-Dihydroxy-12-ursen-3-yl palmitate

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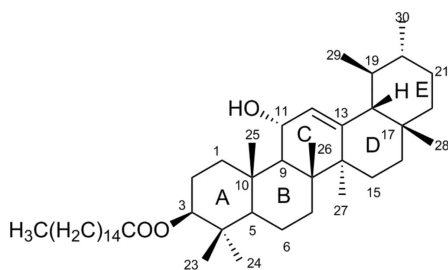
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 Key indicators: single-crystal X-ray study; $T = 93$ K; mean $\sigma(\text{C}-\text{C}) = 0.004$ Å; R factor = 0.043; wR factor = 0.098; data-to-parameter ratio = 10.6.

In the title compound, $\text{C}_{46}\text{H}_{80}\text{O}_3$, a natural ursane-type triperpenoid, four of the five six-membered rings adopt chair conformations; the fifth, which has a $\text{C}=\text{C}$ double bond, adopts an approximate half-boat conformation. In the crystal, molecules are linked by $\text{O}-\text{H}\cdots\text{O}$ hydrogen bonds, forming chains along [010].

Related literature

For standard bond lengths, see: Allen *et al.* (1987). For spectroscopic properties of the title compound, see: Kakuda *et al.* (2003).


Experimental
Crystal data
 $\text{C}_{46}\text{H}_{80}\text{O}_3$
 $M_r = 681.10$

 Monoclinic, $P2_1$
 $a = 11.389$ (2) Å
 $b = 15.714$ (3) Å
 $c = 11.766$ (2) Å
 $\beta = 98.925$ (3)°
 $V = 2080.3$ (7) Å³
 $Z = 2$
 Mo $K\alpha$ radiation
 $\mu = 0.07$ mm⁻¹
 $T = 93$ K
 $0.60 \times 0.50 \times 0.30$ mm

Data collection

 Rigaku AFC10/Saturn724+
 diffractometer
 16995 measured reflections

 4832 independent reflections
 4397 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.032$
Refinement
 $R[F^2 > 2\sigma(F^2)] = 0.043$
 $wR(F^2) = 0.098$
 $S = 1.00$
 4832 reflections
 455 parameters
 1 restraint

 H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\text{max}} = 0.22$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.15$ e Å⁻³
Table 1

Hydrogen-bond geometry (Å, °).

| $D-\text{H}\cdots A$ | $D-\text{H}$ | $\text{H}\cdots A$ | $D\cdots A$ | $D-\text{H}\cdots A$ |
|--|--------------|--------------------|-------------|----------------------|
| $\text{O}1-\text{H}1\text{O}\cdots\text{O}3^i$ | 0.79 (4) | 2.13 (4) | 2.886 (3) | 161 (3) |

 Symmetry code: (i) $-x + 1, y - \frac{1}{2}, -z + 1$.

Data collection: *CrystalClear* (Rigaku, 2008); cell refinement: *CrystalClear*; data reduction: *CrystalClear*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008) and *PLATON* (Spek, 2009); software used to prepare material for publication: *SHELXTL*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: BX2380).

References

- Allen, F. H., Kennard, O., Watson, D. G., Brammer, L., Orpen, A. G. & Taylor, R. (1987). *J. Chem. Soc. Perkin Trans. 2*, pp. S1–19.
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supporting information

Acta Cryst. (2011). E67, o3355 [https://doi.org/10.1107/S1600536811047106]

3 β ,11 α -Dihydroxy-12-ursen-3-yl palmitate**Yong-Liang Yuan, Su-Ping Bai, Hui-Juan Liang and Dan-Dan Ye****S1. Comment**

The title compound, (I), is a naturally occurring ursene-type triterpene isolated from the medicinal plant *Saussurea nivea* Turcz. This plant has been used as antibacterial, inflammation-diminishing drugs and febrifuge. The structure of compound (I) has been reported previously based on spectroscopic methods (Kakuda, *et al.*, 2003). In order to further confirm the structure and conformation of (I), a crystal structure analysis, reported here, was undertaken.

The X-ray crystallographic analysis of (I) confirms the previously proposed molecular structure of (I) as an ursene-type triterpene. Fig. 1 shows its conformation: the palmitoyl group and the hydroxyl group is connected to C3 and C11 in β and α -orientation, respectively; while the Me-20 and Me-30 adopted β and α -orientation at C19 and C20, respectively; and the double bond is located at C12 and C13. The molecule contains five six-membered rings. The A/B and B/C ring junctions show *trans* fusion and the geometry of the rings is *cis* at the D/E ring junction. The bond lengths and angles of (I) have normal values (Allen *et al.*, 1987), with the following average values (Å): $Csp^3-Csp^3 = 1.533$ (3), $Csp^3-Csp^2 = 1.522$ (3), $Csp^2-Csp^2 = 1.333$ (3), $C=O = 1.211$ (3), $Csp^2-O = 1.330$ (3), $Csp^3-O = 1.452$ (3)). Rings A, B and E have slightly flattened chair conformations, with average torsion angles of 53.6 (3), 53.4 (3) and 54.2 °, respectively. Chair conformations of ring D are twisted, with torsion angle range from 34.1 to 63.0 °. Ring C adopts an approximate half-boat conformation. The long carbon chain connected to C3, which takes anti-conformations in C4'-C9' and C10'-C16' with average dihedral angles of 176.9 and 177.0 °, has two turns on C3'-C4' and C9'-C10' in *gauche*-conformations with torsion angles of 73.9 and 69.4°, respectively. The crystal packing is stabilized by intermolecular O—H...O hydrogen bonds involving the carbonyl and hydroxyl groups (Table 1 and Fig. 2). The hydrogen bonds link the molecules into chains along the b axis.

S2. Experimental

The dried and crushed leaves of *Saussurea nivea* Turcz (10 kg, collected from Tongbai Mountain, Henan Province, China) were extracted three times with Me₂CO at room temperature for seven days. The extract was filtered and the solvent was removed under reduced pressure. The residue was partitioned between water and ethylacetate. After removing the solvent, the ethyl acetate residue was separated by repeated silica gel (200–300 mesh) column chromatography and recrystallization in CHCl₃/Me₂CO (30:1) to afford 20 mg of compound (I) (Optical rotation: $[\alpha]_D^{25} +31.1$ ° (c 0.27, CHCl₃). Crystals suitable for x-ray experiment were obtained by slow evaporation of a solution of the compound (I) in Me₂CO at room temperature.

S3. Refinement

All H atoms were included in calculated position and refined as riding atoms, with C—H = 0.95 Å (CH₃), 0.93 and 0.97 Å (CH₂), 0.98 Å (CH), and with $U_{iso}(H) = 1.2U_{eq}(C)$. In the absence of significant anomalous scattering effects, Friedel pairs were merged. The choice of enantiomer was based on comparison of the optical rotation with that of related compounds

with known stereochemistry.

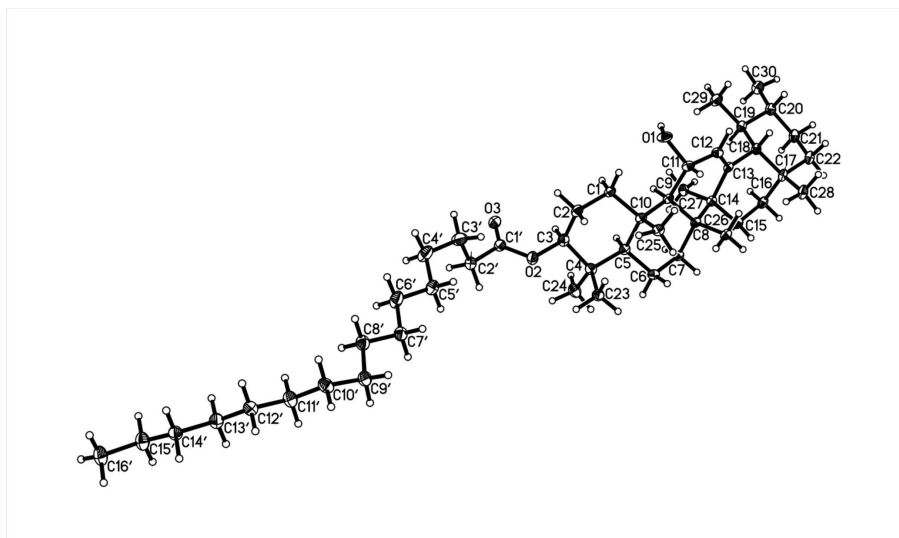


Figure 1

A view of the molecular structure of compound (I). Displacement ellipsoids are drawn at the 50% probability level.

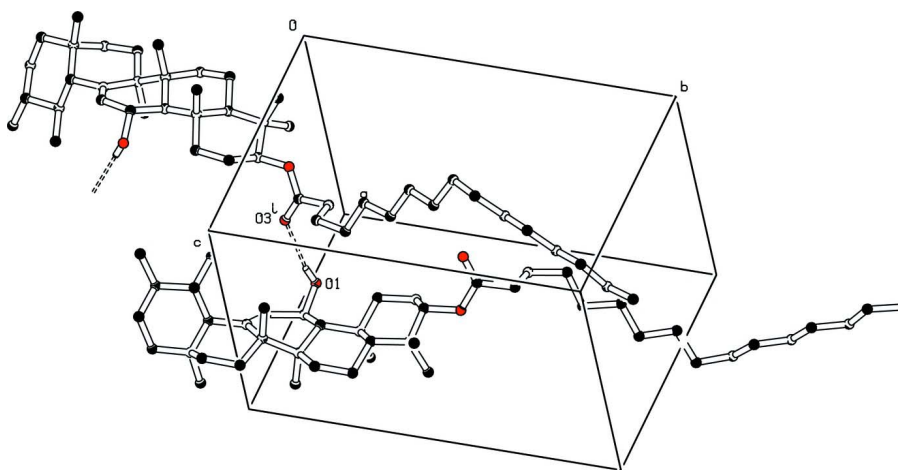


Figure 2

The crystal packing of (I), viewed along the *a* axis, showing the O—H \cdots O hydrogen bonds as dashed lines. Symmetry code: (i) $-x+1, y-1/2, -z+1$.

3 β ,11 α -Dihydroxy-12-ursen-3-yl palmitate

Crystal data

$C_{46}H_{80}O_3$

$M_r = 681.10$

Monoclinic, $P2_1$

$a = 11.389$ (2) Å

$b = 15.714$ (3) Å

$c = 11.766$ (2) Å

$\beta = 98.925$ (3)°

$V = 2080.3$ (7) Å³

$Z = 2$

$F(000) = 760$

$D_x = 1.087$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 7345 reflections

$\theta = 3.0$ – 27.5 °

$\mu = 0.07$ mm⁻¹

$T = 93$ K

Block, colourless

$0.60 \times 0.50 \times 0.30$ mm

Data collection

Rigaku AFC10/Saturn724+
diffractometer

Radiation source: Rotating Anode

Graphite monochromator

Detector resolution: 28.5714 pixels mm⁻¹

phi and ω scans

16995 measured reflections

4832 independent reflections

4397 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.032$

$\theta_{\text{max}} = 27.5^\circ$, $\theta_{\text{min}} = 3.1^\circ$

$h = -14 \rightarrow 14$

$k = -20 \rightarrow 20$

$l = -14 \rightarrow 14$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.043$

$wR(F^2) = 0.098$

$S = 1.00$

4832 reflections

455 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

$w = 1/[\sigma^2(F_o^2) + (0.0286P)^2 + 0.860P]$

where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\text{max}} < 0.001$

$\Delta\rho_{\text{max}} = 0.22 \text{ e } \text{\AA}^{-3}$

$\Delta\rho_{\text{min}} = -0.15 \text{ e } \text{\AA}^{-3}$

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)

| | <i>x</i> | <i>y</i> | <i>z</i> | $U_{\text{iso}}^*/U_{\text{eq}}$ |
|-----|--------------|--------------|--------------|----------------------------------|
| O1 | 0.7358 (2) | 0.09757 (13) | 0.56526 (16) | 0.0423 (5) |
| O2 | 0.66023 (16) | 0.51467 (12) | 0.64282 (16) | 0.0373 (4) |
| O3 | 0.48079 (18) | 0.51257 (14) | 0.53336 (17) | 0.0469 (5) |
| C1 | 0.7086 (2) | 0.27759 (15) | 0.6092 (2) | 0.0282 (5) |
| H1A | 0.7494 | 0.2468 | 0.5531 | 0.034* |
| H1B | 0.6248 | 0.2584 | 0.5980 | 0.034* |
| C2 | 0.7121 (2) | 0.37373 (16) | 0.5849 (2) | 0.0321 (6) |
| H2A | 0.7958 | 0.3925 | 0.5903 | 0.039* |
| H2B | 0.6716 | 0.3853 | 0.5057 | 0.039* |
| C3 | 0.6522 (2) | 0.42345 (17) | 0.6692 (2) | 0.0328 (6) |
| H3 | 0.5664 | 0.4069 | 0.6590 | 0.039* |
| C4 | 0.7061 (2) | 0.41063 (16) | 0.7958 (2) | 0.0302 (5) |
| C5 | 0.7115 (2) | 0.31251 (16) | 0.8183 (2) | 0.0263 (5) |
| H5 | 0.6264 | 0.2944 | 0.8085 | 0.032* |
| C6 | 0.7608 (2) | 0.28945 (16) | 0.9424 (2) | 0.0298 (5) |
| H6A | 0.7300 | 0.3299 | 0.9952 | 0.036* |
| H6B | 0.8485 | 0.2940 | 0.9545 | 0.036* |

| | | | | |
|------|--------------|---------------|--------------|------------|
| C7 | 0.7248 (2) | 0.19890 (16) | 0.9696 (2) | 0.0286 (5) |
| H7A | 0.6374 | 0.1967 | 0.9651 | 0.034* |
| H7B | 0.7605 | 0.1850 | 1.0496 | 0.034* |
| C8 | 0.76315 (19) | 0.13074 (16) | 0.8885 (2) | 0.0245 (5) |
| C9 | 0.73602 (19) | 0.15932 (16) | 0.76015 (19) | 0.0238 (5) |
| H9 | 0.6476 | 0.1558 | 0.7394 | 0.029* |
| C10 | 0.7685 (2) | 0.25440 (16) | 0.7329 (2) | 0.0253 (5) |
| C11 | 0.7853 (2) | 0.09126 (17) | 0.6844 (2) | 0.0288 (5) |
| H11 | 0.8727 | 0.1016 | 0.6903 | 0.035* |
| C12 | 0.7700 (2) | 0.00159 (16) | 0.7251 (2) | 0.0274 (5) |
| H12 | 0.7920 | -0.0427 | 0.6775 | 0.033* |
| C13 | 0.72899 (19) | -0.02268 (15) | 0.8202 (2) | 0.0241 (5) |
| C14 | 0.6930 (2) | 0.04465 (16) | 0.9029 (2) | 0.0251 (5) |
| C15 | 0.7157 (2) | 0.01434 (17) | 1.0297 (2) | 0.0307 (5) |
| H15A | 0.7993 | 0.0270 | 1.0626 | 0.037* |
| H15B | 0.6641 | 0.0473 | 1.0740 | 0.037* |
| C16 | 0.6926 (2) | -0.08099 (17) | 1.0454 (2) | 0.0318 (5) |
| H16A | 0.6064 | -0.0922 | 1.0249 | 0.038* |
| H16B | 0.7164 | -0.0962 | 1.1273 | 0.038* |
| C17 | 0.7605 (2) | -0.13792 (17) | 0.9712 (2) | 0.0293 (5) |
| C18 | 0.7145 (2) | -0.11755 (16) | 0.8435 (2) | 0.0268 (5) |
| H18 | 0.7681 | -0.1483 | 0.7976 | 0.032* |
| C19 | 0.5867 (2) | -0.15270 (16) | 0.8029 (2) | 0.0288 (5) |
| H19 | 0.5315 | -0.1230 | 0.8482 | 0.035* |
| C20 | 0.5810 (2) | -0.24905 (17) | 0.8281 (2) | 0.0320 (5) |
| H20 | 0.6385 | -0.2786 | 0.7854 | 0.038* |
| C21 | 0.6193 (2) | -0.26572 (17) | 0.9567 (2) | 0.0350 (6) |
| H21A | 0.5630 | -0.2375 | 1.0008 | 0.042* |
| H21B | 0.6170 | -0.3276 | 0.9717 | 0.042* |
| C22 | 0.7438 (2) | -0.23238 (18) | 0.9970 (2) | 0.0351 (6) |
| H22A | 0.8006 | -0.2661 | 0.9596 | 0.042* |
| H22B | 0.7638 | -0.2414 | 1.0810 | 0.042* |
| C23 | 0.8264 (2) | 0.45627 (17) | 0.8259 (2) | 0.0326 (6) |
| H23A | 0.8791 | 0.4389 | 0.7716 | 0.039* |
| H23B | 0.8628 | 0.4410 | 0.9042 | 0.039* |
| H23C | 0.8142 | 0.5180 | 0.8209 | 0.039* |
| C24 | 0.6193 (2) | 0.45184 (18) | 0.8686 (3) | 0.0413 (7) |
| H24A | 0.6090 | 0.5122 | 0.8485 | 0.050* |
| H24B | 0.6516 | 0.4465 | 0.9505 | 0.050* |
| H24C | 0.5422 | 0.4229 | 0.8531 | 0.050* |
| C25 | 0.9039 (2) | 0.26877 (17) | 0.7378 (2) | 0.0289 (5) |
| H25A | 0.9380 | 0.2211 | 0.7003 | 0.035* |
| H25B | 0.9420 | 0.2725 | 0.8182 | 0.035* |
| H25C | 0.9172 | 0.3218 | 0.6979 | 0.035* |
| C26 | 0.8995 (2) | 0.11656 (17) | 0.9255 (2) | 0.0294 (5) |
| H26A | 0.9419 | 0.1698 | 0.9165 | 0.035* |
| H26B | 0.9271 | 0.0725 | 0.8770 | 0.035* |
| H26C | 0.9150 | 0.0985 | 1.0062 | 0.035* |

| | | | | |
|------|------------|---------------|------------|------------|
| C27 | 0.5556 (2) | 0.05609 (17) | 0.8704 (2) | 0.0296 (5) |
| H27A | 0.5349 | 0.0629 | 0.7869 | 0.036* |
| H27B | 0.5304 | 0.1067 | 0.9090 | 0.036* |
| H27C | 0.5152 | 0.0058 | 0.8949 | 0.036* |
| C28 | 0.8948 (2) | -0.11911 (19) | 0.9984 (2) | 0.0394 (6) |
| H28A | 0.9383 | -0.1596 | 0.9570 | 0.047* |
| H28B | 0.9215 | -0.1248 | 1.0813 | 0.047* |
| H28C | 0.9101 | -0.0610 | 0.9741 | 0.047* |
| C29 | 0.5451 (3) | -0.13413 (19) | 0.6761 (2) | 0.0426 (7) |
| H29A | 0.5984 | -0.1620 | 0.6298 | 0.051* |
| H29B | 0.5458 | -0.0725 | 0.6631 | 0.051* |
| H29C | 0.4642 | -0.1559 | 0.6537 | 0.051* |
| C30 | 0.4576 (2) | -0.28749 (19) | 0.7892 (2) | 0.0398 (6) |
| H30A | 0.4577 | -0.3473 | 0.8127 | 0.048* |
| H30B | 0.4379 | -0.2836 | 0.7053 | 0.048* |
| H30C | 0.3985 | -0.2561 | 0.8248 | 0.048* |
| C1' | 0.5716 (2) | 0.54980 (18) | 0.5713 (2) | 0.0350 (6) |
| C2' | 0.5993 (3) | 0.64035 (18) | 0.5409 (2) | 0.0390 (6) |
| H2'1 | 0.5272 | 0.6759 | 0.5379 | 0.047* |
| H2'2 | 0.6615 | 0.6639 | 0.6007 | 0.047* |
| C3' | 0.6418 (3) | 0.6424 (2) | 0.4257 (3) | 0.0492 (8) |
| H3'1 | 0.5834 | 0.6122 | 0.3687 | 0.059* |
| H3'2 | 0.7181 | 0.6113 | 0.4320 | 0.059* |
| C4' | 0.6596 (3) | 0.7331 (2) | 0.3812 (3) | 0.0549 (9) |
| H4'1 | 0.6651 | 0.7296 | 0.2982 | 0.066* |
| H4'2 | 0.5883 | 0.7672 | 0.3891 | 0.066* |
| C5' | 0.7677 (3) | 0.77948 (19) | 0.4411 (2) | 0.0403 (6) |
| H5'1 | 0.8392 | 0.7444 | 0.4376 | 0.048* |
| H5'2 | 0.7601 | 0.7874 | 0.5231 | 0.048* |
| C6' | 0.7836 (3) | 0.8663 (2) | 0.3869 (3) | 0.0511 (8) |
| H6'1 | 0.7864 | 0.8577 | 0.3040 | 0.061* |
| H6'2 | 0.7125 | 0.9012 | 0.3930 | 0.061* |
| C7' | 0.8925 (2) | 0.91628 (18) | 0.4381 (2) | 0.0369 (6) |
| H7'1 | 0.9642 | 0.8808 | 0.4367 | 0.044* |
| H7'2 | 0.8875 | 0.9296 | 0.5195 | 0.044* |
| C8' | 0.9052 (3) | 0.99922 (19) | 0.3730 (3) | 0.0407 (6) |
| H8'1 | 0.9087 | 0.9854 | 0.2915 | 0.049* |
| H8'2 | 0.8333 | 1.0342 | 0.3750 | 0.049* |
| C9' | 1.0140 (3) | 1.05190 (18) | 0.4201 (3) | 0.0413 (7) |
| H9'1 | 1.0856 | 1.0156 | 0.4244 | 0.050* |
| H9'2 | 1.0071 | 1.0704 | 0.4992 | 0.050* |
| C10' | 1.0300 (2) | 1.13012 (18) | 0.3475 (3) | 0.0414 (7) |
| H10A | 1.0274 | 1.1123 | 0.2665 | 0.050* |
| H10B | 1.1096 | 1.1547 | 0.3740 | 0.050* |
| C11' | 0.9365 (3) | 1.19878 (19) | 0.3527 (3) | 0.0424 (7) |
| H11A | 0.9364 | 1.2146 | 0.4341 | 0.051* |
| H11B | 0.8573 | 1.1750 | 0.3225 | 0.051* |
| C12' | 0.9563 (2) | 1.27902 (18) | 0.2846 (2) | 0.0382 (6) |

| | | | | |
|-------|------------|--------------|------------|-------------|
| H12A | 1.0346 | 1.3038 | 0.3156 | 0.046* |
| H12B | 0.9574 | 1.2635 | 0.2032 | 0.046* |
| C13' | 0.8604 (2) | 1.34523 (19) | 0.2901 (3) | 0.0396 (6) |
| H13A | 0.8564 | 1.3576 | 0.3719 | 0.048* |
| H13B | 0.7829 | 1.3208 | 0.2556 | 0.048* |
| C14' | 0.8783 (2) | 1.42849 (18) | 0.2295 (2) | 0.0357 (6) |
| H14A | 0.9517 | 1.4562 | 0.2686 | 0.043* |
| H14B | 0.8894 | 1.4162 | 0.1493 | 0.043* |
| C15' | 0.7755 (3) | 1.4889 (2) | 0.2281 (3) | 0.0464 (7) |
| H15C | 0.7619 | 1.4983 | 0.3082 | 0.056* |
| H15D | 0.7032 | 1.4618 | 0.1857 | 0.056* |
| C16' | 0.7929 (3) | 1.5745 (2) | 0.1735 (3) | 0.0502 (8) |
| H16'A | 0.8613 | 1.6035 | 0.2178 | 0.060* |
| H16'B | 0.7215 | 1.6093 | 0.1733 | 0.060* |
| H16'C | 0.8072 | 1.5660 | 0.0943 | 0.060* |
| H1O | 0.674 (3) | 0.074 (3) | 0.554 (3) | 0.057 (11)* |

Atomic displacement parameters (Å²)

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|-----|-------------|-------------|-------------|--------------|--------------|--------------|
| O1 | 0.0667 (14) | 0.0380 (11) | 0.0214 (10) | -0.0177 (11) | 0.0045 (9) | -0.0005 (8) |
| O2 | 0.0324 (9) | 0.0323 (10) | 0.0443 (11) | -0.0016 (8) | -0.0027 (8) | 0.0066 (9) |
| O3 | 0.0447 (11) | 0.0427 (11) | 0.0469 (12) | 0.0035 (10) | -0.0134 (9) | -0.0033 (10) |
| C1 | 0.0303 (12) | 0.0304 (13) | 0.0224 (12) | -0.0072 (10) | -0.0010 (9) | 0.0018 (10) |
| C2 | 0.0325 (12) | 0.0337 (14) | 0.0284 (13) | -0.0080 (11) | -0.0006 (10) | 0.0063 (11) |
| C3 | 0.0245 (12) | 0.0316 (13) | 0.0411 (15) | -0.0042 (10) | 0.0009 (10) | 0.0036 (12) |
| C4 | 0.0253 (12) | 0.0293 (13) | 0.0364 (14) | -0.0021 (10) | 0.0057 (10) | -0.0010 (11) |
| C5 | 0.0221 (11) | 0.0303 (12) | 0.0267 (13) | -0.0026 (9) | 0.0039 (9) | 0.0001 (10) |
| C6 | 0.0312 (12) | 0.0323 (13) | 0.0261 (13) | -0.0021 (10) | 0.0054 (10) | -0.0045 (10) |
| C7 | 0.0304 (12) | 0.0346 (14) | 0.0208 (12) | -0.0002 (10) | 0.0044 (9) | 0.0002 (10) |
| C8 | 0.0205 (10) | 0.0308 (12) | 0.0219 (12) | -0.0010 (9) | 0.0026 (8) | 0.0005 (10) |
| C9 | 0.0204 (10) | 0.0305 (12) | 0.0196 (11) | -0.0041 (9) | 0.0005 (8) | 0.0008 (9) |
| C10 | 0.0224 (11) | 0.0291 (12) | 0.0242 (12) | -0.0053 (9) | 0.0031 (9) | 0.0001 (10) |
| C11 | 0.0304 (12) | 0.0333 (13) | 0.0231 (12) | -0.0075 (10) | 0.0057 (10) | -0.0007 (10) |
| C12 | 0.0264 (11) | 0.0304 (13) | 0.0262 (12) | -0.0026 (10) | 0.0067 (9) | -0.0030 (10) |
| C13 | 0.0199 (10) | 0.0300 (12) | 0.0214 (12) | 0.0007 (9) | 0.0003 (9) | 0.0020 (9) |
| C14 | 0.0224 (11) | 0.0310 (12) | 0.0223 (12) | 0.0010 (9) | 0.0047 (9) | 0.0019 (9) |
| C15 | 0.0321 (12) | 0.0385 (14) | 0.0218 (12) | 0.0015 (11) | 0.0053 (10) | 0.0030 (11) |
| C16 | 0.0321 (12) | 0.0404 (14) | 0.0224 (12) | 0.0015 (11) | 0.0027 (10) | 0.0089 (11) |
| C17 | 0.0235 (11) | 0.0346 (13) | 0.0285 (13) | 0.0035 (10) | 0.0000 (9) | 0.0103 (11) |
| C18 | 0.0240 (11) | 0.0299 (13) | 0.0271 (13) | 0.0036 (9) | 0.0059 (9) | 0.0046 (10) |
| C19 | 0.0283 (12) | 0.0307 (13) | 0.0271 (13) | 0.0011 (10) | 0.0031 (9) | 0.0049 (10) |
| C20 | 0.0317 (13) | 0.0320 (13) | 0.0329 (14) | -0.0004 (10) | 0.0071 (10) | 0.0066 (11) |
| C21 | 0.0369 (14) | 0.0334 (14) | 0.0355 (15) | 0.0008 (11) | 0.0081 (11) | 0.0118 (11) |
| C22 | 0.0320 (13) | 0.0382 (15) | 0.0341 (14) | 0.0056 (11) | 0.0024 (11) | 0.0119 (12) |
| C23 | 0.0313 (13) | 0.0309 (13) | 0.0349 (14) | -0.0039 (10) | 0.0028 (10) | -0.0002 (11) |
| C24 | 0.0387 (15) | 0.0358 (15) | 0.0519 (18) | 0.0037 (12) | 0.0152 (13) | 0.0013 (13) |
| C25 | 0.0236 (11) | 0.0329 (13) | 0.0306 (13) | -0.0055 (10) | 0.0052 (9) | 0.0002 (10) |

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|------|-------------|-------------|-------------|--------------|--------------|--------------|
| C26 | 0.0235 (11) | 0.0376 (14) | 0.0256 (12) | -0.0009 (10) | -0.0006 (9) | 0.0008 (10) |
| C27 | 0.0217 (11) | 0.0323 (13) | 0.0357 (14) | 0.0027 (10) | 0.0072 (10) | 0.0030 (11) |
| C28 | 0.0277 (13) | 0.0424 (16) | 0.0453 (16) | 0.0058 (12) | -0.0034 (11) | 0.0110 (13) |
| C29 | 0.0504 (16) | 0.0360 (15) | 0.0366 (15) | -0.0131 (13) | -0.0082 (12) | 0.0079 (12) |
| C30 | 0.0389 (15) | 0.0412 (16) | 0.0398 (16) | -0.0052 (12) | 0.0074 (12) | 0.0088 (13) |
| C1' | 0.0362 (13) | 0.0364 (14) | 0.0313 (14) | 0.0044 (11) | 0.0018 (11) | -0.0012 (11) |
| C2' | 0.0428 (15) | 0.0351 (15) | 0.0373 (15) | 0.0054 (12) | 0.0005 (12) | 0.0027 (12) |
| C3' | 0.0579 (19) | 0.0502 (19) | 0.0373 (17) | -0.0116 (15) | 0.0007 (13) | -0.0043 (14) |
| C4' | 0.057 (2) | 0.066 (2) | 0.0379 (17) | -0.0130 (17) | -0.0059 (14) | 0.0150 (16) |
| C5' | 0.0404 (15) | 0.0463 (17) | 0.0338 (15) | 0.0011 (12) | 0.0046 (11) | 0.0078 (13) |
| C6' | 0.0489 (17) | 0.054 (2) | 0.0472 (18) | -0.0057 (15) | -0.0046 (14) | 0.0187 (15) |
| C7' | 0.0400 (14) | 0.0384 (15) | 0.0336 (14) | 0.0072 (12) | 0.0104 (11) | 0.0007 (12) |
| C8' | 0.0456 (15) | 0.0388 (15) | 0.0388 (15) | 0.0046 (13) | 0.0101 (12) | 0.0040 (12) |
| C9' | 0.0404 (15) | 0.0389 (15) | 0.0439 (17) | 0.0098 (12) | 0.0046 (12) | -0.0022 (13) |
| C10' | 0.0367 (14) | 0.0356 (15) | 0.0534 (18) | 0.0005 (12) | 0.0115 (13) | -0.0058 (14) |
| C11' | 0.0398 (15) | 0.0373 (15) | 0.0516 (18) | 0.0040 (12) | 0.0114 (13) | 0.0041 (13) |
| C12' | 0.0379 (14) | 0.0349 (15) | 0.0420 (16) | -0.0005 (11) | 0.0075 (12) | -0.0010 (12) |
| C13' | 0.0332 (14) | 0.0396 (15) | 0.0454 (16) | -0.0024 (12) | 0.0042 (12) | 0.0044 (13) |
| C14' | 0.0322 (13) | 0.0344 (14) | 0.0399 (15) | -0.0026 (11) | 0.0036 (11) | 0.0013 (12) |
| C15' | 0.0385 (15) | 0.0436 (17) | 0.0577 (19) | 0.0051 (13) | 0.0095 (14) | 0.0114 (14) |
| C16' | 0.0411 (16) | 0.0417 (17) | 0.068 (2) | 0.0077 (13) | 0.0100 (15) | 0.0155 (16) |

Geometric parameters (Å, °)

| | | | |
|--------|-----------|----------|-----------|
| O1—C11 | 1.431 (3) | C24—H24B | 0.9800 |
| O1—H1O | 0.79 (4) | C24—H24C | 0.9800 |
| O2—C1' | 1.330 (3) | C25—H25A | 0.9800 |
| O2—C3 | 1.473 (3) | C25—H25B | 0.9800 |
| O3—C1' | 1.211 (3) | C25—H25C | 0.9800 |
| C1—C2 | 1.539 (3) | C26—H26A | 0.9800 |
| C1—C10 | 1.552 (3) | C26—H26B | 0.9800 |
| C1—H1A | 0.9900 | C26—H26C | 0.9800 |
| C1—H1B | 0.9900 | C27—H27A | 0.9800 |
| C2—C3 | 1.507 (4) | C27—H27B | 0.9800 |
| C2—H2A | 0.9900 | C27—H27C | 0.9800 |
| C2—H2B | 0.9900 | C28—H28A | 0.9800 |
| C3—C4 | 1.533 (4) | C28—H28B | 0.9800 |
| C3—H3 | 1.0000 | C28—H28C | 0.9800 |
| C4—C23 | 1.539 (3) | C29—H29A | 0.9800 |
| C4—C24 | 1.547 (4) | C29—H29B | 0.9800 |
| C4—C5 | 1.564 (3) | C29—H29C | 0.9800 |
| C5—C6 | 1.524 (3) | C30—H30A | 0.9800 |
| C5—C10 | 1.571 (3) | C30—H30B | 0.9800 |
| C5—H5 | 1.0000 | C30—H30C | 0.9800 |
| C6—C7 | 1.528 (3) | C1'—C2' | 1.513 (4) |
| C6—H6A | 0.9900 | C2'—C3' | 1.508 (4) |
| C6—H6B | 0.9900 | C2'—H2'1 | 0.9900 |
| C7—C8 | 1.542 (3) | C2'—H2'2 | 0.9900 |

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|------------|-----------|---------------|-----------|
| C7—H7A | 0.9900 | C3'—C4' | 1.542 (5) |
| C7—H7B | 0.9900 | C3'—H3'1 | 0.9900 |
| C8—C9 | 1.559 (3) | C3'—H3'2 | 0.9900 |
| C8—C26 | 1.562 (3) | C4'—C5' | 1.508 (4) |
| C8—C14 | 1.594 (3) | C4'—H4'1 | 0.9900 |
| C9—C11 | 1.552 (3) | C4'—H4'2 | 0.9900 |
| C9—C10 | 1.584 (3) | C5'—C6' | 1.528 (4) |
| C9—H9 | 1.0000 | C5'—H5'1 | 0.9900 |
| C10—C25 | 1.550 (3) | C5'—H5'2 | 0.9900 |
| C11—C12 | 1.507 (4) | C6'—C7' | 1.512 (4) |
| C11—H11 | 1.0000 | C6'—H6'1 | 0.9900 |
| C12—C13 | 1.333 (3) | C6'—H6'2 | 0.9900 |
| C12—H12 | 0.9500 | C7'—C8' | 1.530 (4) |
| C13—C18 | 1.530 (3) | C7'—H7'1 | 0.9900 |
| C13—C14 | 1.536 (3) | C7'—H7'2 | 0.9900 |
| C14—C15 | 1.549 (3) | C8'—C9' | 1.521 (4) |
| C14—C27 | 1.563 (3) | C8'—H8'1 | 0.9900 |
| C15—C16 | 1.537 (4) | C8'—H8'2 | 0.9900 |
| C15—H15A | 0.9900 | C9'—C10' | 1.524 (4) |
| C15—H15B | 0.9900 | C9'—H9'1 | 0.9900 |
| C16—C17 | 1.539 (4) | C9'—H9'2 | 0.9900 |
| C16—H16A | 0.9900 | C10'—C11' | 1.524 (4) |
| C16—H16B | 0.9900 | C10'—H10A | 0.9900 |
| C17—C22 | 1.533 (4) | C10'—H10B | 0.9900 |
| C17—C28 | 1.542 (3) | C11'—C12' | 1.529 (4) |
| C17—C18 | 1.546 (3) | C11'—H11A | 0.9900 |
| C18—C19 | 1.561 (3) | C11'—H11B | 0.9900 |
| C18—H18 | 1.0000 | C12'—C13' | 1.517 (4) |
| C19—C29 | 1.522 (3) | C12'—H12A | 0.9900 |
| C19—C20 | 1.546 (4) | C12'—H12B | 0.9900 |
| C19—H19 | 1.0000 | C13'—C14' | 1.519 (4) |
| C20—C21 | 1.531 (4) | C13'—H13A | 0.9900 |
| C20—C30 | 1.533 (4) | C13'—H13B | 0.9900 |
| C20—H20 | 1.0000 | C14'—C15' | 1.505 (4) |
| C21—C22 | 1.517 (4) | C14'—H14A | 0.9900 |
| C21—H21A | 0.9900 | C14'—H14B | 0.9900 |
| C21—H21B | 0.9900 | C15'—C16' | 1.517 (4) |
| C22—H22A | 0.9900 | C15'—H15C | 0.9900 |
| C22—H22B | 0.9900 | C15'—H15D | 0.9900 |
| C23—H23A | 0.9800 | C16'—H16'A | 0.9800 |
| C23—H23B | 0.9800 | C16'—H16'B | 0.9800 |
| C23—H23C | 0.9800 | C16'—H16'C | 0.9800 |
| C24—H24A | 0.9800 | | |
| C11—O1—H1O | 110 (3) | C4—C24—H24B | 109.5 |
| C1'—O2—C3 | 118.1 (2) | H24A—C24—H24B | 109.5 |
| C2—C1—C10 | 112.6 (2) | C4—C24—H24C | 109.5 |
| C2—C1—H1A | 109.1 | H24A—C24—H24C | 109.5 |

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| C10—C1—H1A | 109.1 | H24B—C24—H24C | 109.5 |
| C2—C1—H1B | 109.1 | C10—C25—H25A | 109.5 |
| C10—C1—H1B | 109.1 | C10—C25—H25B | 109.5 |
| H1A—C1—H1B | 107.8 | H25A—C25—H25B | 109.5 |
| C3—C2—C1 | 111.0 (2) | C10—C25—H25C | 109.5 |
| C3—C2—H2A | 109.4 | H25A—C25—H25C | 109.5 |
| C1—C2—H2A | 109.4 | H25B—C25—H25C | 109.5 |
| C3—C2—H2B | 109.4 | C8—C26—H26A | 109.5 |
| C1—C2—H2B | 109.4 | C8—C26—H26B | 109.5 |
| H2A—C2—H2B | 108.0 | H26A—C26—H26B | 109.5 |
| O2—C3—C2 | 108.4 (2) | C8—C26—H26C | 109.5 |
| O2—C3—C4 | 107.7 (2) | H26A—C26—H26C | 109.5 |
| C2—C3—C4 | 114.7 (2) | H26B—C26—H26C | 109.5 |
| O2—C3—H3 | 108.6 | C14—C27—H27A | 109.5 |
| C2—C3—H3 | 108.6 | C14—C27—H27B | 109.5 |
| C4—C3—H3 | 108.6 | H27A—C27—H27B | 109.5 |
| C3—C4—C23 | 111.7 (2) | C14—C27—H27C | 109.5 |
| C3—C4—C24 | 106.8 (2) | H27A—C27—H27C | 109.5 |
| C23—C4—C24 | 107.5 (2) | H27B—C27—H27C | 109.5 |
| C3—C4—C5 | 107.1 (2) | C17—C28—H28A | 109.5 |
| C23—C4—C5 | 114.2 (2) | C17—C28—H28B | 109.5 |
| C24—C4—C5 | 109.3 (2) | H28A—C28—H28B | 109.5 |
| C6—C5—C4 | 113.3 (2) | C17—C28—H28C | 109.5 |
| C6—C5—C10 | 110.3 (2) | H28A—C28—H28C | 109.5 |
| C4—C5—C10 | 118.3 (2) | H28B—C28—H28C | 109.5 |
| C6—C5—H5 | 104.4 | C19—C29—H29A | 109.5 |
| C4—C5—H5 | 104.4 | C19—C29—H29B | 109.5 |
| C10—C5—H5 | 104.4 | H29A—C29—H29B | 109.5 |
| C5—C6—C7 | 110.6 (2) | C19—C29—H29C | 109.5 |
| C5—C6—H6A | 109.5 | H29A—C29—H29C | 109.5 |
| C7—C6—H6A | 109.5 | H29B—C29—H29C | 109.5 |
| C5—C6—H6B | 109.5 | C20—C30—H30A | 109.5 |
| C7—C6—H6B | 109.5 | C20—C30—H30B | 109.5 |
| H6A—C6—H6B | 108.1 | H30A—C30—H30B | 109.5 |
| C6—C7—C8 | 113.97 (19) | C20—C30—H30C | 109.5 |
| C6—C7—H7A | 108.8 | H30A—C30—H30C | 109.5 |
| C8—C7—H7A | 108.8 | H30B—C30—H30C | 109.5 |
| C6—C7—H7B | 108.8 | O3—C1'—O2 | 123.6 (3) |
| C8—C7—H7B | 108.8 | O3—C1'—C2' | 124.4 (3) |
| H7A—C7—H7B | 107.7 | O2—C1'—C2' | 112.0 (2) |
| C7—C8—C9 | 111.5 (2) | C3'—C2'—C1' | 109.9 (2) |
| C7—C8—C26 | 106.91 (19) | C3'—C2'—H2'1 | 109.7 |
| C9—C8—C26 | 110.50 (18) | C1'—C2'—H2'1 | 109.7 |
| C7—C8—C14 | 109.20 (18) | C3'—C2'—H2'2 | 109.7 |
| C9—C8—C14 | 108.69 (18) | C1'—C2'—H2'2 | 109.7 |
| C26—C8—C14 | 110.02 (19) | H2'1—C2'—H2'2 | 108.2 |
| C11—C9—C8 | 108.81 (19) | C2'—C3'—C4' | 113.7 (3) |
| C11—C9—C10 | 114.66 (18) | C2'—C3'—H3'1 | 108.8 |

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| C8—C9—C10 | 116.76 (18) | C4'—C3'—H3'1 | 108.8 |
| C11—C9—H9 | 105.1 | C2'—C3'—H3'2 | 108.8 |
| C8—C9—H9 | 105.1 | C4'—C3'—H3'2 | 108.8 |
| C10—C9—H9 | 105.1 | H3'1—C3'—H3'2 | 107.7 |
| C25—C10—C1 | 106.87 (19) | C5'—C4'—C3' | 115.4 (3) |
| C25—C10—C5 | 113.51 (19) | C5'—C4'—H4'1 | 108.4 |
| C1—C10—C5 | 107.37 (19) | C3'—C4'—H4'1 | 108.4 |
| C25—C10—C9 | 113.08 (19) | C5'—C4'—H4'2 | 108.4 |
| C1—C10—C9 | 109.20 (18) | C3'—C4'—H4'2 | 108.4 |
| C5—C10—C9 | 106.63 (18) | H4'1—C4'—H4'2 | 107.5 |
| O1—C11—C12 | 109.2 (2) | C4'—C5'—C6' | 112.2 (2) |
| O1—C11—C9 | 113.1 (2) | C4'—C5'—H5'1 | 109.2 |
| C12—C11—C9 | 113.08 (19) | C6'—C5'—H5'1 | 109.2 |
| O1—C11—H11 | 107.0 | C4'—C5'—H5'2 | 109.2 |
| C12—C11—H11 | 107.0 | C6'—C5'—H5'2 | 109.2 |
| C9—C11—H11 | 107.0 | H5'1—C5'—H5'2 | 107.9 |
| C13—C12—C11 | 127.4 (2) | C7'—C6'—C5' | 116.1 (2) |
| C13—C12—H12 | 116.3 | C7'—C6'—H6'1 | 108.3 |
| C11—C12—H12 | 116.3 | C5'—C6'—H6'1 | 108.3 |
| C12—C13—C18 | 119.4 (2) | C7'—C6'—H6'2 | 108.3 |
| C12—C13—C14 | 119.8 (2) | C5'—C6'—H6'2 | 108.3 |
| C18—C13—C14 | 120.69 (19) | H6'1—C6'—H6'2 | 107.4 |
| C13—C14—C15 | 112.2 (2) | C6'—C7'—C8' | 112.2 (2) |
| C13—C14—C27 | 106.24 (19) | C6'—C7'—H7'1 | 109.2 |
| C15—C14—C27 | 106.42 (19) | C8'—C7'—H7'1 | 109.2 |
| C13—C14—C8 | 109.24 (18) | C6'—C7'—H7'2 | 109.2 |
| C15—C14—C8 | 110.61 (19) | C8'—C7'—H7'2 | 109.2 |
| C27—C14—C8 | 112.09 (19) | H7'1—C7'—H7'2 | 107.9 |
| C16—C15—C14 | 114.1 (2) | C9'—C8'—C7' | 114.6 (2) |
| C16—C15—H15A | 108.7 | C9'—C8'—H8'1 | 108.6 |
| C14—C15—H15A | 108.7 | C7'—C8'—H8'1 | 108.6 |
| C16—C15—H15B | 108.7 | C9'—C8'—H8'2 | 108.6 |
| C14—C15—H15B | 108.7 | C7'—C8'—H8'2 | 108.6 |
| H15A—C15—H15B | 107.6 | H8'1—C8'—H8'2 | 107.6 |
| C15—C16—C17 | 112.8 (2) | C8'—C9'—C10' | 113.4 (2) |
| C15—C16—H16A | 109.0 | C8'—C9'—H9'1 | 108.9 |
| C17—C16—H16A | 109.0 | C10'—C9'—H9'1 | 108.9 |
| C15—C16—H16B | 109.0 | C8'—C9'—H9'2 | 108.9 |
| C17—C16—H16B | 109.0 | C10'—C9'—H9'2 | 108.9 |
| H16A—C16—H16B | 107.8 | H9'1—C9'—H9'2 | 107.7 |
| C22—C17—C16 | 111.3 (2) | C9'—C10'—C11' | 113.8 (2) |
| C22—C17—C28 | 107.1 (2) | C9'—C10'—H10A | 108.8 |
| C16—C17—C28 | 109.9 (2) | C11'—C10'—H10A | 108.8 |
| C22—C17—C18 | 111.0 (2) | C9'—C10'—H10B | 108.8 |
| C16—C17—C18 | 107.82 (19) | C11'—C10'—H10B | 108.8 |
| C28—C17—C18 | 109.7 (2) | H10A—C10'—H10B | 107.7 |
| C13—C18—C17 | 110.3 (2) | C10'—C11'—C12' | 113.7 (2) |
| C13—C18—C19 | 114.29 (19) | C10'—C11'—H11A | 108.8 |

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| C17—C18—C19 | 112.55 (19) | C12'—C11'—H11A | 108.8 |
| C13—C18—H18 | 106.4 | C10'—C11'—H11B | 108.8 |
| C17—C18—H18 | 106.4 | C12'—C11'—H11B | 108.8 |
| C19—C18—H18 | 106.4 | H11A—C11'—H11B | 107.7 |
| C29—C19—C20 | 111.1 (2) | C13'—C12'—C11' | 112.2 (2) |
| C29—C19—C18 | 111.5 (2) | C13'—C12'—H12A | 109.2 |
| C20—C19—C18 | 110.7 (2) | C11'—C12'—H12A | 109.2 |
| C29—C19—H19 | 107.8 | C13'—C12'—H12B | 109.2 |
| C20—C19—H19 | 107.8 | C11'—C12'—H12B | 109.2 |
| C18—C19—H19 | 107.8 | H12A—C12'—H12B | 107.9 |
| C21—C20—C30 | 109.6 (2) | C12'—C13'—C14' | 114.9 (2) |
| C21—C20—C19 | 110.1 (2) | C12'—C13'—H13A | 108.5 |
| C30—C20—C19 | 113.1 (2) | C14'—C13'—H13A | 108.5 |
| C21—C20—H20 | 108.0 | C12'—C13'—H13B | 108.5 |
| C30—C20—H20 | 108.0 | C14'—C13'—H13B | 108.5 |
| C19—C20—H20 | 108.0 | H13A—C13'—H13B | 107.5 |
| C22—C21—C20 | 111.0 (2) | C15'—C14'—C13' | 112.8 (2) |
| C22—C21—H21A | 109.4 | C15'—C14'—H14A | 109.0 |
| C20—C21—H21A | 109.4 | C13'—C14'—H14A | 109.0 |
| C22—C21—H21B | 109.4 | C15'—C14'—H14B | 109.0 |
| C20—C21—H21B | 109.4 | C13'—C14'—H14B | 109.0 |
| H21A—C21—H21B | 108.0 | H14A—C14'—H14B | 107.8 |
| C21—C22—C17 | 114.3 (2) | C14'—C15'—C16' | 114.3 (2) |
| C21—C22—H22A | 108.7 | C14'—C15'—H15C | 108.7 |
| C17—C22—H22A | 108.7 | C16'—C15'—H15C | 108.7 |
| C21—C22—H22B | 108.7 | C14'—C15'—H15D | 108.7 |
| C17—C22—H22B | 108.7 | C16'—C15'—H15D | 108.7 |
| H22A—C22—H22B | 107.6 | H15C—C15'—H15D | 107.6 |
| C4—C23—H23A | 109.5 | C15'—C16'—H16'A | 109.5 |
| C4—C23—H23B | 109.5 | C15'—C16'—H16'B | 109.5 |
| H23A—C23—H23B | 109.5 | H16'A—C16'—H16'B | 109.5 |
| C4—C23—H23C | 109.5 | C15'—C16'—H16'C | 109.5 |
| H23A—C23—H23C | 109.5 | H16'A—C16'—H16'C | 109.5 |
| H23B—C23—H23C | 109.5 | H16'B—C16'—H16'C | 109.5 |
| C4—C24—H24A | 109.5 | | |
| | | | |
| C10—C1—C2—C3 | -58.1 (3) | C7—C8—C14—C13 | 178.74 (18) |
| C1'—O2—C3—C2 | 91.8 (3) | C9—C8—C14—C13 | 56.9 (2) |
| C1'—O2—C3—C4 | -143.6 (2) | C26—C8—C14—C13 | -64.2 (2) |
| C1—C2—C3—O2 | 179.05 (19) | C7—C8—C14—C15 | -57.3 (2) |
| C1—C2—C3—C4 | 58.7 (3) | C9—C8—C14—C15 | -179.17 (19) |
| O2—C3—C4—C23 | -47.3 (3) | C26—C8—C14—C15 | 59.7 (2) |
| C2—C3—C4—C23 | 73.5 (3) | C7—C8—C14—C27 | 61.3 (2) |
| O2—C3—C4—C24 | 70.0 (2) | C9—C8—C14—C27 | -60.6 (2) |
| C2—C3—C4—C24 | -169.2 (2) | C26—C8—C14—C27 | 178.31 (19) |
| O2—C3—C4—C5 | -173.05 (19) | C13—C14—C15—C16 | -36.2 (3) |
| C2—C3—C4—C5 | -52.2 (3) | C27—C14—C15—C16 | 79.6 (2) |
| C3—C4—C5—C6 | -178.62 (19) | C8—C14—C15—C16 | -158.42 (19) |

| | | | |
|-----------------|--------------|---------------------|-------------|
| C23—C4—C5—C6 | 57.2 (3) | C14—C15—C16—C17 | 53.3 (3) |
| C24—C4—C5—C6 | -63.3 (3) | C15—C16—C17—C22 | 175.0 (2) |
| C3—C4—C5—C10 | 50.0 (3) | C15—C16—C17—C28 | 56.6 (3) |
| C23—C4—C5—C10 | -74.2 (3) | C15—C16—C17—C18 | -63.0 (3) |
| C24—C4—C5—C10 | 165.3 (2) | C12—C13—C18—C17 | 136.7 (2) |
| C4—C5—C6—C7 | 160.3 (2) | C14—C13—C18—C17 | -45.3 (3) |
| C10—C5—C6—C7 | -64.5 (2) | C12—C13—C18—C19 | -95.3 (3) |
| C5—C6—C7—C8 | 56.4 (3) | C14—C13—C18—C19 | 82.7 (3) |
| C6—C7—C8—C9 | -45.0 (3) | C22—C17—C18—C13 | 178.59 (19) |
| C6—C7—C8—C26 | 75.8 (2) | C16—C17—C18—C13 | 56.5 (2) |
| C6—C7—C8—C14 | -165.16 (19) | C28—C17—C18—C13 | -63.2 (3) |
| C7—C8—C9—C11 | 175.71 (19) | C22—C17—C18—C19 | 49.6 (3) |
| C26—C8—C9—C11 | 57.0 (2) | C16—C17—C18—C19 | -72.5 (3) |
| C14—C8—C9—C11 | -63.9 (2) | C28—C17—C18—C19 | 167.8 (2) |
| C7—C8—C9—C10 | 44.0 (2) | C13—C18—C19—C29 | 54.9 (3) |
| C26—C8—C9—C10 | -74.8 (2) | C17—C18—C19—C29 | -178.3 (2) |
| C14—C8—C9—C10 | 164.42 (17) | C13—C18—C19—C20 | 179.0 (2) |
| C2—C1—C10—C25 | -70.0 (3) | C17—C18—C19—C20 | -54.1 (3) |
| C2—C1—C10—C5 | 52.1 (3) | C29—C19—C20—C21 | -178.1 (2) |
| C2—C1—C10—C9 | 167.35 (19) | C18—C19—C20—C21 | 57.5 (3) |
| C6—C5—C10—C25 | -65.6 (3) | C29—C19—C20—C30 | -55.2 (3) |
| C4—C5—C10—C25 | 67.2 (3) | C18—C19—C20—C30 | -179.5 (2) |
| C6—C5—C10—C1 | 176.55 (19) | C30—C20—C21—C22 | 177.1 (2) |
| C4—C5—C10—C1 | -50.7 (3) | C19—C20—C21—C22 | -57.9 (3) |
| C6—C5—C10—C9 | 59.6 (2) | C20—C21—C22—C17 | 55.4 (3) |
| C4—C5—C10—C9 | -167.66 (19) | C16—C17—C22—C21 | 69.4 (3) |
| C11—C9—C10—C25 | -54.3 (3) | C28—C17—C22—C21 | -170.5 (2) |
| C8—C9—C10—C25 | 74.7 (2) | C18—C17—C22—C21 | -50.7 (3) |
| C11—C9—C10—C1 | 64.6 (2) | C3—O2—C1'—O3 | 5.6 (4) |
| C8—C9—C10—C1 | -166.46 (19) | C3—O2—C1'—C2' | -172.8 (2) |
| C11—C9—C10—C5 | -179.70 (19) | O3—C1'—C2'—C3' | -79.7 (3) |
| C8—C9—C10—C5 | -50.7 (2) | O2—C1'—C2'—C3' | 98.7 (3) |
| C8—C9—C11—O1 | 162.55 (19) | C1'—C2'—C3'—C4' | 173.8 (3) |
| C10—C9—C11—O1 | -64.6 (2) | C2'—C3'—C4'—C5' | 73.9 (4) |
| C8—C9—C11—C12 | 37.8 (3) | C3'—C4'—C5'—C6' | 176.2 (3) |
| C10—C9—C11—C12 | 170.67 (18) | C4'—C5'—C6'—C7' | -177.6 (3) |
| O1—C11—C12—C13 | -132.5 (3) | C5'—C6'—C7'—C8' | 176.1 (3) |
| C9—C11—C12—C13 | -5.7 (3) | C6'—C7'—C8'—C9' | -179.5 (3) |
| C11—C12—C13—C18 | 177.1 (2) | C7'—C8'—C9'—C10' | 175.0 (2) |
| C11—C12—C13—C14 | -0.9 (4) | C8'—C9'—C10'—C11' | 69.4 (3) |
| C12—C13—C14—C15 | -147.9 (2) | C9'—C10'—C11'—C12' | 177.2 (2) |
| C18—C13—C14—C15 | 34.1 (3) | C10'—C11'—C12'—C13' | 179.0 (3) |
| C12—C13—C14—C27 | 96.2 (2) | C11'—C12'—C13'—C14' | 176.9 (2) |
| C18—C13—C14—C27 | -81.7 (2) | C12'—C13'—C14'—C15' | 174.7 (3) |
| C12—C13—C14—C8 | -24.9 (3) | C13'—C14'—C15'—C16' | 177.1 (3) |
| C18—C13—C14—C8 | 157.2 (2) | | |

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—H1O···O3 ⁱ | 0.79 (4) | 2.13 (4) | 2.886 (3) | 161 (3) |

Symmetry code: (i) $-x+1, y-1/2, -z+1$.