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3-Hydroxy-1,2-dimethoxyanthraquinone

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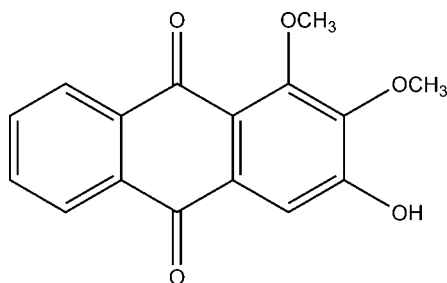
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Key indicators: single-crystal X-ray study; $T = 293$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; R factor = 0.044; wR factor = 0.136; data-to-parameter ratio = 10.4.

The title compound, $\text{C}_{16}\text{H}_{12}\text{O}_5$, was isolated from *Morinda officinalis* How. The anthraquinone ring system is almost planar, the dihedral angle between the two benzene rings being 1.12 (4°). In the crystal structure, $\text{O}-\text{H}\cdots\text{O}$ and $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds link the molecules in the crystallographic a -axis direction. Weak $\pi-\pi$ stacking interactions [centroid-centroid distance between symmetry-related benzene rings of 3.699 (4) Å] are also present.

Related literature

For the biological properties of anthraquinone derivatives, see: Kim *et al.* (2005) and of the title compound, see: Ali *et al.* (2000); Jia *et al.* (2007); Wu *et al.* (2003). For related structures, see: Ng *et al.* (2005); Boonnak *et al.* (2005). For the structure of another compound isolated from *Morinda officinalis* How., see: Liu & Jiao (2009). For reference structural data, see: Allen *et al.* (1987).



Experimental

Crystal data

$\text{C}_{16}\text{H}_{12}\text{O}_5$
 $M_r = 284.26$
Triclinic, $P\bar{1}$
 $a = 7.4087$ (17) Å

$b = 8.0387$ (18) Å
 $c = 11.802$ (3) Å
 $\alpha = 95.386$ (3°)
 $\beta = 92.357$ (3°)

$\gamma = 115.712$ (2°)
 $V = 627.9$ (3) Å³
 $Z = 2$
Mo $K\alpha$ radiation

$\mu = 0.11$ mm⁻¹
 $T = 293$ K
 $0.30 \times 0.20 \times 0.20$ mm

Data collection

Bruker APEXII area-detector diffractometer
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.967$, $T_{\max} = 0.978$

3200 measured reflections
2182 independent reflections
1639 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.013$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.044$
 $wR(F^2) = 0.136$
 $S = 1.06$
2182 reflections
210 parameters

H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\max} = 0.25$ e Å⁻³
 $\Delta\rho_{\min} = -0.17$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

| $D-H\cdots A$ | $D-H$ | $H\cdots A$ | $D\cdots A$ | $D-H\cdots A$ |
|---|----------|-------------|-------------|---------------|
| $\text{C}25-\text{H}5\cdots\text{O}1^i$ | 0.97 (3) | 2.57 (3) | 3.256 (2) | 128 (2) |
| $\text{O}8-\text{H}6\cdots\text{O}1^i$ | 0.88 (3) | 1.91 (3) | 2.781 (2) | 168 (3) |

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

Data collection: APEX2 (Bruker, 2004); cell refinement: SAINT (Bruker, 2004); data reduction: SAINT (Bruker, 2004); program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: XP in SHELXTL (Sheldrick, 2008); 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: PK2164).

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supporting information

Acta Cryst. (2009). E65, o1524 [doi:10.1107/S1600536809021266]

3-Hydroxy-1,2-dimethoxyanthraquinone

Yong-Jun Xu, Xiao-Xi Yang and Hong-Bin Zhao

S1. Comment

Anthraquinone derivatives extracted from the roots of *Morinda officinalis* How. (most common familiar name in China: Bajitian) have been used to support the entire body treating a wide range of symptoms, including poor digestion, high blood pressure and immune deficiencies in China since ancient times. Recent studies have demonstrated that they have multiple pharmacological actions such as anti-HIV, anti-inflammatory, antinociceptive, antimicrobial, antioxidant, antihepatotoxic and antimutagenic activities (Kim et al., 2005). One component found in *Morinda officinalis* How., 1,2-dimethoxy-3-hydroxyanthraquinone, exhibits a variety of potent biological effects such as antiviral and antimicrobial activities (Ali et al., 2000), antioxidant activity (Jia et al., 2007) and cytotoxic activity (Wu et al., 2003). We report here the structure of the title compound.

In the title compound (Fig. 1), the C-C bond lengths show normal values (Allen et al., 1987), and the C-O and C=O bond lengths are comparable to those observed in similar structures (Ng et al., 2005; Boonnak et al., 2005). The anthraquinone ring system is substantially planar, the dihedral angle between the two benzene rings being $1.12(4)^\circ$. The molecules are self-assembled by C—H \cdots O and O—H \cdots O hydrogen bonding interactions (Table 1) into a supramolecular network. The crystal structure is further stabilized by weak π - π interactions along the a axis (Fig. 2) between the anthraquinone ring systems of the stacked molecules. The centroid-to-centroid distances between related benzene rings of the stacked molecules is $3.699(4)\text{\AA}$, thus indicating weak π - π contacts.

S2. Experimental

The roots of *Morinda officinalis* How. (1000 g) were shattered to powder (about 30 mesh) and extracted with 85% ethanol (4000 ml) for 2 h with stirring. The extraction procedure was repeated three times. The extracts were combined and evaporated to dryness under reduced pressure at 333 K, the residue was redissolved in water (800 ml). Then the enriched extracts were extracted with chloroform three times (800 ml each), the chloroform solutions were combined and evaporated to dryness under reduced pressure at 333 K, 6.80 g crude extracts were obtained. The crude extracts were separated with n-hexane-ethyl acetate-methanol-water (6 : 4 : 5 : 5, v/v) using high-speed counter-current chromatography (HSCCC) to obtain 1,2-dimethoxy-3-hydroxyanthraquinone (yield 20.3 mg). Single crystals suitable for X-ray analysis were obtained by slow evaporation of a methanol solution.

S3. Refinement

Methyl H atoms were placed at calculated positions and treated as riding on the parent C atoms with C—H = 0.96 \AA and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$. Coordinates of all other hydrogens were refined but their U_{iso} values were fixed at 0.105 \AA^2 .

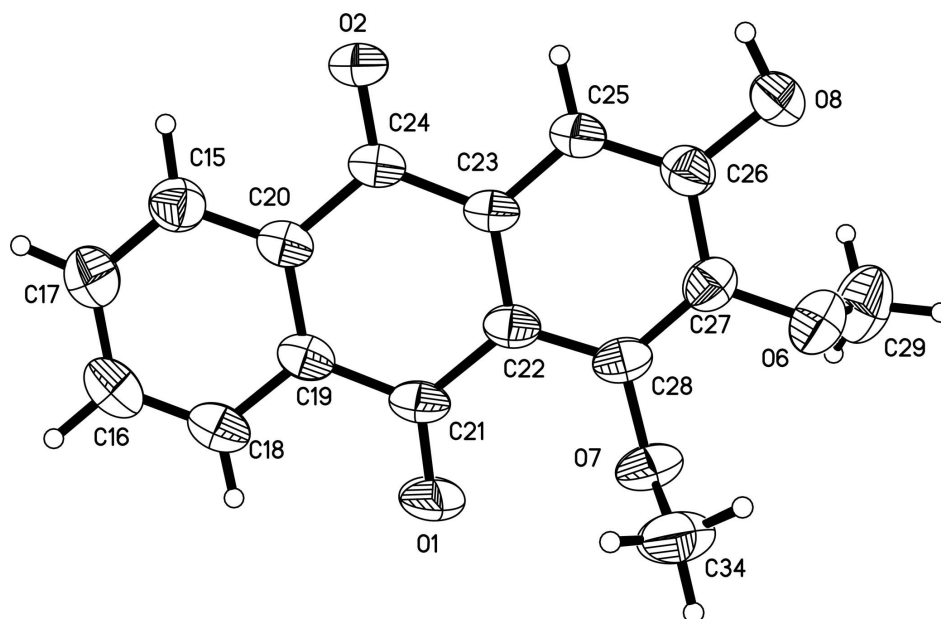


Figure 1

The molecular structure showing the atomic-numbering scheme and displacement ellipsoids drawn at the 30% probability level.

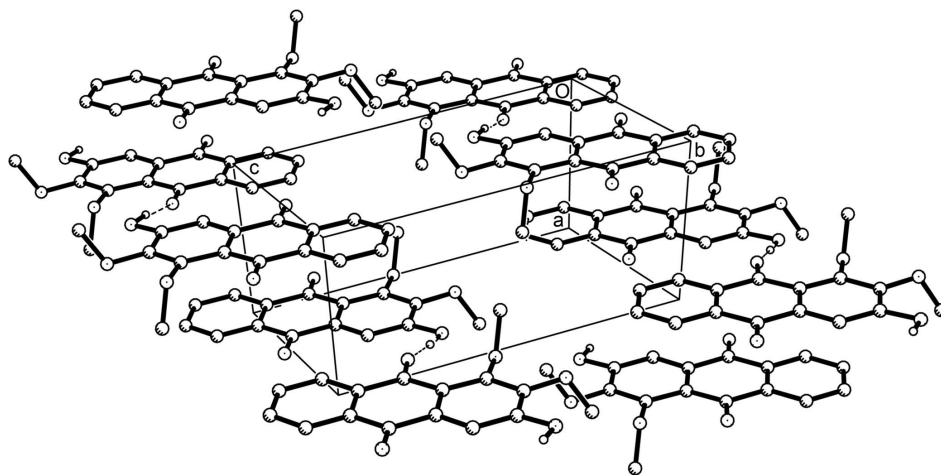


Figure 2

The molecular packing showing the hydrogen bonding interactions as broken lines.

3-Hydroxy-1,2-dimethoxyanthraquinone

Crystal data

$C_{16}H_{12}O_5$

$M_r = 284.26$

Triclinic, $P\bar{1}$

Hall symbol: $-P\ 1$

$a = 7.4087\ (17)\ \text{\AA}$

$b = 8.0387\ (18)\ \text{\AA}$

$c = 11.802\ (3)\ \text{\AA}$

$\alpha = 95.386\ (3)^\circ$

$\beta = 92.357\ (3)^\circ$

$\gamma = 115.712\ (2)^\circ$

$V = 627.9\ (3)\ \text{\AA}^3$

$Z = 2$

$F(000) = 296.0$

$D_x = 1.503\ \text{Mg m}^{-3}$

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

Cell parameters from 1305 reflections

$\theta = 3.1\text{--}27.3^\circ$
 $\mu = 0.11 \text{ mm}^{-1}$
 $T = 293 \text{ K}$

Block, yellow
 $0.30 \times 0.20 \times 0.20 \text{ mm}$

Data collection

Bruker APEXII area-detector
 diffractometer
 Radiation source: fine-focus sealed tube
 Graphite monochromator
 φ and ω scans
 Absorption correction: multi-scan
 (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.967$, $T_{\max} = 0.978$

3200 measured reflections
 2182 independent reflections
 1639 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.013$
 $\theta_{\max} = 25.0^\circ$, $\theta_{\min} = 2.8^\circ$
 $h = -7 \rightarrow 8$
 $k = -9 \rightarrow 9$
 $l = -14 \rightarrow 11$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.044$
 $wR(F^2) = 0.136$
 $S = 1.06$
 2182 reflections
 210 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 atoms treated by a mixture of independent
 and constrained refinement
 $w = 1/[\sigma^2(F_o^2) + (0.0736P)^2 + 0.1017P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.25 \text{ e } \text{Å}^{-3}$
 $\Delta\rho_{\min} = -0.17 \text{ e } \text{Å}^{-3}$

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 (Å^2)

| | <i>x</i> | <i>y</i> | <i>z</i> | $U_{\text{iso}}^*/U_{\text{eq}}$ |
|-----|------------|--------------|---------------|----------------------------------|
| O1 | 0.2605 (2) | 0.79647 (18) | 0.14638 (13) | 0.0605 (4) |
| O2 | 0.2359 (2) | 0.18299 (17) | -0.09353 (11) | 0.0560 (4) |
| O6 | 0.2197 (2) | 0.3418 (2) | 0.43860 (11) | 0.0597 (4) |
| O7 | 0.2427 (2) | 0.64028 (18) | 0.34133 (12) | 0.0547 (4) |
| O8 | 0.1946 (2) | 0.03307 (19) | 0.30888 (12) | 0.0582 (4) |
| C15 | 0.2626 (3) | 0.4756 (3) | -0.21242 (16) | 0.0463 (5) |
| C16 | 0.2985 (3) | 0.7870 (3) | -0.20248 (19) | 0.0550 (5) |
| C17 | 0.2839 (3) | 0.6275 (3) | -0.26601 (19) | 0.0546 (5) |
| C18 | 0.2898 (3) | 0.7953 (3) | -0.08620 (19) | 0.0474 (5) |
| C19 | 0.2686 (2) | 0.6434 (2) | -0.03041 (15) | 0.0379 (4) |
| C20 | 0.2551 (2) | 0.4817 (2) | -0.09503 (15) | 0.0372 (4) |
| C21 | 0.2574 (3) | 0.6545 (2) | 0.09545 (16) | 0.0403 (4) |

| | | | | |
|------|------------|------------|---------------|------------|
| C22 | 0.2422 (2) | 0.4946 (2) | 0.15295 (15) | 0.0368 (4) |
| C23 | 0.2284 (2) | 0.3315 (2) | 0.08725 (14) | 0.0350 (4) |
| C24 | 0.2378 (2) | 0.3203 (2) | -0.03826 (15) | 0.0377 (4) |
| C25 | 0.2095 (3) | 0.1777 (2) | 0.13812 (16) | 0.0399 (4) |
| C26 | 0.2058 (3) | 0.1783 (2) | 0.25489 (15) | 0.0425 (4) |
| C27 | 0.2176 (3) | 0.3354 (3) | 0.32218 (15) | 0.0440 (5) |
| C28 | 0.2364 (3) | 0.4919 (2) | 0.27130 (15) | 0.0405 (4) |
| C29 | 0.0304 (4) | 0.2459 (4) | 0.4794 (2) | 0.0804 (8) |
| H29A | -0.0439 | 0.3180 | 0.4754 | 0.097* |
| H29B | 0.0483 | 0.2264 | 0.5573 | 0.097* |
| H29C | -0.0422 | 0.1278 | 0.4333 | 0.097* |
| C34 | 0.4369 (4) | 0.7632 (3) | 0.3940 (2) | 0.0703 (7) |
| H34A | 0.4862 | 0.6969 | 0.4403 | 0.084* |
| H34B | 0.4295 | 0.8636 | 0.4412 | 0.084* |
| H34C | 0.5266 | 0.8121 | 0.3363 | 0.084* |
| H1 | 0.249 (4) | 0.360 (4) | -0.255 (2) | 0.105* |
| H4 | 0.298 (5) | 0.897 (4) | -0.044 (3) | 0.105* |
| H3 | 0.323 (4) | 0.897 (4) | -0.239 (2) | 0.105* |
| H2 | 0.292 (4) | 0.616 (4) | -0.345 (3) | 0.105* |
| H5 | 0.198 (4) | 0.069 (4) | 0.089 (3) | 0.105* |
| H6 | 0.199 (5) | -0.044 (4) | 0.253 (3) | 0.105* |

Atomic displacement parameters (\AA^2)

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|-----|-------------|-------------|-------------|-------------|--------------|--------------|
| O1 | 0.0852 (11) | 0.0373 (7) | 0.0687 (9) | 0.0365 (7) | 0.0148 (8) | -0.0003 (6) |
| O2 | 0.0889 (11) | 0.0383 (7) | 0.0479 (8) | 0.0358 (7) | 0.0051 (7) | -0.0014 (6) |
| O6 | 0.0681 (10) | 0.0683 (10) | 0.0419 (8) | 0.0301 (8) | 0.0044 (7) | 0.0023 (7) |
| O7 | 0.0608 (9) | 0.0478 (8) | 0.0564 (8) | 0.0291 (7) | 0.0023 (7) | -0.0142 (6) |
| O8 | 0.0884 (11) | 0.0506 (9) | 0.0507 (8) | 0.0423 (8) | 0.0151 (7) | 0.0147 (6) |
| C15 | 0.0462 (11) | 0.0455 (11) | 0.0489 (11) | 0.0220 (9) | 0.0019 (8) | 0.0054 (8) |
| C16 | 0.0524 (12) | 0.0451 (12) | 0.0689 (14) | 0.0202 (10) | 0.0041 (10) | 0.0208 (10) |
| C17 | 0.0569 (13) | 0.0551 (12) | 0.0534 (12) | 0.0247 (10) | 0.0048 (10) | 0.0143 (10) |
| C18 | 0.0420 (11) | 0.0346 (10) | 0.0673 (13) | 0.0183 (9) | 0.0037 (9) | 0.0076 (9) |
| C19 | 0.0293 (9) | 0.0313 (9) | 0.0540 (11) | 0.0144 (7) | 0.0026 (7) | 0.0046 (8) |
| C20 | 0.0295 (9) | 0.0320 (9) | 0.0501 (11) | 0.0140 (7) | 0.0020 (7) | 0.0032 (7) |
| C21 | 0.0349 (9) | 0.0291 (9) | 0.0575 (11) | 0.0161 (8) | 0.0050 (8) | -0.0014 (8) |
| C22 | 0.0312 (9) | 0.0307 (9) | 0.0491 (11) | 0.0152 (7) | 0.0045 (7) | 0.0000 (7) |
| C23 | 0.0310 (9) | 0.0276 (9) | 0.0460 (10) | 0.0135 (7) | 0.0034 (7) | -0.0005 (7) |
| C24 | 0.0347 (9) | 0.0289 (9) | 0.0480 (10) | 0.0140 (7) | 0.0012 (7) | -0.0004 (7) |
| C25 | 0.0419 (10) | 0.0316 (9) | 0.0476 (11) | 0.0179 (8) | 0.0055 (8) | 0.0020 (7) |
| C26 | 0.0451 (11) | 0.0388 (10) | 0.0472 (11) | 0.0213 (9) | 0.0067 (8) | 0.0071 (8) |
| C27 | 0.0439 (11) | 0.0496 (11) | 0.0400 (10) | 0.0228 (9) | 0.0043 (8) | 0.0008 (8) |
| C28 | 0.0363 (10) | 0.0386 (10) | 0.0470 (10) | 0.0191 (8) | 0.0027 (7) | -0.0060 (8) |
| C29 | 0.0866 (18) | 0.102 (2) | 0.0514 (13) | 0.0379 (16) | 0.0221 (12) | 0.0158 (13) |
| C34 | 0.0775 (16) | 0.0484 (12) | 0.0745 (15) | 0.0243 (12) | -0.0103 (12) | -0.0172 (11) |

Geometric parameters (Å, °)

| | | | |
|-------------|-------------|---------------|-------------|
| O1—C21 | 1.230 (2) | C19—C21 | 1.488 (3) |
| O2—C24 | 1.222 (2) | C20—C24 | 1.475 (2) |
| O6—C27 | 1.369 (2) | C21—C22 | 1.473 (3) |
| O6—C29 | 1.407 (3) | C22—C28 | 1.401 (3) |
| O7—C28 | 1.368 (2) | C22—C23 | 1.421 (2) |
| O7—C34 | 1.418 (3) | C23—C25 | 1.380 (3) |
| O8—C26 | 1.355 (2) | C23—C24 | 1.482 (3) |
| O8—H6 | 0.88 (3) | C25—C26 | 1.379 (3) |
| C15—C17 | 1.379 (3) | C25—H5 | 0.97 (3) |
| C15—C20 | 1.386 (3) | C26—C27 | 1.394 (3) |
| C15—H1 | 0.98 (3) | C27—C28 | 1.400 (3) |
| C16—C18 | 1.373 (3) | C29—H29A | 0.9600 |
| C16—C17 | 1.382 (3) | C29—H29B | 0.9600 |
| C16—H3 | 0.97 (3) | C29—H29C | 0.9600 |
| C17—H2 | 0.94 (3) | C34—H34A | 0.9600 |
| C18—C19 | 1.394 (3) | C34—H34B | 0.9600 |
| C18—H4 | 0.90 (3) | C34—H34C | 0.9600 |
| C19—C20 | 1.405 (2) | | |
| C27—O6—C29 | 114.93 (16) | C22—C23—C24 | 121.57 (16) |
| C28—O7—C34 | 114.31 (15) | O2—C24—C20 | 120.49 (16) |
| C26—O8—H6 | 102 (2) | O2—C24—C23 | 121.23 (16) |
| C17—C15—C20 | 120.31 (19) | C20—C24—C23 | 118.26 (15) |
| C17—C15—H1 | 122.3 (18) | C26—C25—C23 | 120.99 (16) |
| C20—C15—H1 | 117.3 (18) | C26—C25—H5 | 121.2 (18) |
| C18—C16—C17 | 120.59 (19) | C23—C25—H5 | 117.8 (18) |
| C18—C16—H3 | 119.1 (17) | O8—C26—C25 | 123.02 (16) |
| C17—C16—H3 | 120.2 (17) | O8—C26—C27 | 117.56 (16) |
| C15—C17—C16 | 120.0 (2) | C25—C26—C27 | 119.41 (17) |
| C15—C17—H2 | 116.1 (19) | O6—C27—C26 | 120.69 (17) |
| C16—C17—H2 | 123.9 (19) | O6—C27—C28 | 119.27 (16) |
| C16—C18—C19 | 120.33 (19) | C26—C27—C28 | 119.98 (17) |
| C16—C18—H4 | 122 (2) | O7—C28—C27 | 117.34 (16) |
| C19—C18—H4 | 118 (2) | O7—C28—C22 | 121.20 (17) |
| C18—C19—C20 | 119.00 (18) | C27—C28—C22 | 121.44 (15) |
| C18—C19—C21 | 119.49 (16) | O6—C29—H29A | 109.5 |
| C20—C19—C21 | 121.50 (16) | O6—C29—H29B | 109.5 |
| C15—C20—C19 | 119.80 (16) | H29A—C29—H29B | 109.5 |
| C15—C20—C24 | 119.84 (16) | O6—C29—H29C | 109.5 |
| C19—C20—C24 | 120.34 (16) | H29A—C29—H29C | 109.5 |
| O1—C21—C22 | 123.18 (18) | H29B—C29—H29C | 109.5 |
| O1—C21—C19 | 118.37 (17) | O7—C34—H34A | 109.5 |
| C22—C21—C19 | 118.45 (14) | O7—C34—H34B | 109.5 |
| C28—C22—C23 | 116.89 (16) | H34A—C34—H34B | 109.5 |
| C28—C22—C21 | 123.33 (15) | O7—C34—H34C | 109.5 |
| C23—C22—C21 | 119.77 (16) | H34A—C34—H34C | 109.5 |

| | | | |
|-----------------|--------------|-----------------|--------------|
| C25—C23—C22 | 121.27 (17) | H34B—C34—H34C | 109.5 |
| C25—C23—C24 | 117.15 (15) | | |
| C20—C15—C17—C16 | 0.1 (3) | C19—C20—C24—C23 | 2.0 (2) |
| C18—C16—C17—C15 | -0.6 (3) | C25—C23—C24—O2 | -2.5 (3) |
| C17—C16—C18—C19 | 0.7 (3) | C22—C23—C24—O2 | 176.29 (16) |
| C16—C18—C19—C20 | -0.4 (3) | C25—C23—C24—C20 | 179.16 (14) |
| C16—C18—C19—C21 | -179.35 (16) | C22—C23—C24—C20 | -2.1 (2) |
| C17—C15—C20—C19 | 0.2 (3) | C22—C23—C25—C26 | -0.8 (3) |
| C17—C15—C20—C24 | -178.33 (16) | C24—C23—C25—C26 | 178.01 (15) |
| C18—C19—C20—C15 | -0.1 (3) | C23—C25—C26—O8 | -177.61 (16) |
| C21—C19—C20—C15 | 178.84 (15) | C23—C25—C26—C27 | 1.2 (3) |
| C18—C19—C20—C24 | 178.43 (15) | C29—O6—C27—C26 | -78.9 (2) |
| C21—C19—C20—C24 | -2.6 (2) | C29—O6—C27—C28 | 103.9 (2) |
| C18—C19—C21—O1 | 2.4 (3) | O8—C26—C27—O6 | 0.7 (3) |
| C20—C19—C21—O1 | -176.58 (15) | C25—C26—C27—O6 | -178.22 (16) |
| C18—C19—C21—C22 | -177.85 (15) | O8—C26—C27—C28 | 177.81 (16) |
| C20—C19—C21—C22 | 3.2 (2) | C25—C26—C27—C28 | -1.1 (3) |
| O1—C21—C22—C28 | -2.1 (3) | C34—O7—C28—C27 | 85.1 (2) |
| C19—C21—C22—C28 | 178.18 (15) | C34—O7—C28—C22 | -96.8 (2) |
| O1—C21—C22—C23 | 176.54 (16) | O6—C27—C28—O7 | -4.2 (3) |
| C19—C21—C22—C23 | -3.2 (2) | C26—C27—C28—O7 | 178.56 (15) |
| C28—C22—C23—C25 | 0.2 (3) | O6—C27—C28—C22 | 177.70 (15) |
| C21—C22—C23—C25 | -178.54 (15) | C26—C27—C28—C22 | 0.5 (3) |
| C28—C22—C23—C24 | -178.54 (14) | C23—C22—C28—O7 | -178.03 (14) |
| C21—C22—C23—C24 | 2.8 (2) | C21—C22—C28—O7 | 0.6 (3) |
| C15—C20—C24—O2 | 2.2 (3) | C23—C22—C28—C27 | 0.0 (3) |
| C19—C20—C24—O2 | -176.41 (16) | C21—C22—C28—C27 | 178.61 (15) |
| C15—C20—C24—C23 | -179.46 (15) | | |

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

| $D-H\cdots A$ | $D-H$ | $H\cdots A$ | $D\cdots A$ | $D-H\cdots A$ |
|---------------------------------|----------|-------------|-------------|---------------|
| C25—H5 \cdots O1 ⁱ | 0.97 (3) | 2.57 (3) | 3.256 (2) | 128 (2) |
| O8—H6 \cdots O1 ⁱ | 0.88 (3) | 1.91 (3) | 2.781 (2) | 168 (3) |

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