

Crystal structure of cholest-5-en-3 β -yl 3-(2,4-dimethoxy-3-methylphenyl)prop-2-enoate

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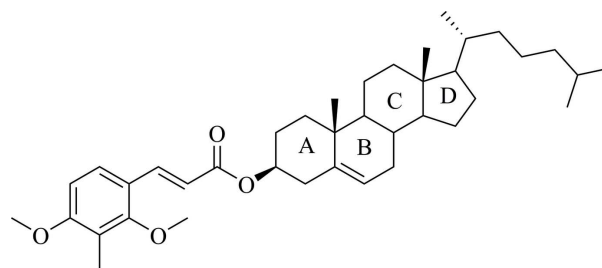
In the title compound, C₃₉H₅₈O₄, the steroid rings *A* and *C* adopt a chair conformation, while ring *B* adopts a half-chair conformation, and ring *D* has an envelope conformation, with the methyl-substituted *C* atom as the flap. In the crystal, molecules pack within layers parallel to (100), with their long axis parallel to the [101] direction. Adjacent layers are linked via C—H \cdots O hydrogen bonds and C—H \cdots π interactions, forming a three-dimensional framework.

Keywords: crystal structure; cholesteryl cinnamates; methyl (*E*)-3-(2,4-dimethoxy-3-methylphenyl)acrylate; hydrogen bonding; C—H \cdots π interactions.

CCDC reference: 884045

1. Related literature

For the preparation of the title compound, see: Thiemann *et al.* (2011). For applications of cholesteryl cinnamates, see: Vora (1976); Kutulya *et al.* (1983); Tanaka *et al.* (1981); Dong *et al.* (2010). For the crystal structure of a similar compound, see: Bugenhagen *et al.* (2012).



2. Experimental

2.1. Crystal data

| | |
|--|-----------------------------------|
| C ₃₉ H ₅₈ O ₄ | $V = 3437.2$ (4) Å ³ |
| $M_r = 590.85$ | $Z = 4$ |
| Orthorhombic, $P2_12_12_1$ | Mo $K\alpha$ radiation |
| $a = 9.4626$ (6) Å | $\mu = 0.07$ mm ⁻¹ |
| $b = 12.2687$ (8) Å | $T = 100$ K |
| $c = 29.6074$ (18) Å | $0.25 \times 0.07 \times 0.07$ mm |

2.2. Data collection

| | |
|--|--|
| Bruker SMART APEX CCD area-detector diffractometer | 33550 measured reflections |
| Absorption correction: multi-scan (SADABS; Bruker, 2009) | 4399 independent reflections |
| $T_{\min} = 0.702$, $T_{\max} = 0.746$ | 3563 reflections with $I > 2\sigma(I)$ |
| | $R_{\text{int}} = 0.052$ |

2.3. Refinement

| | |
|---------------------------------|---|
| $R[F^2 > 2\sigma(F^2)] = 0.046$ | 396 parameters |
| $wR(F^2) = 0.112$ | H-atom parameters constrained |
| $S = 1.03$ | $\Delta\rho_{\text{max}} = 0.56$ e Å ⁻³ |
| 4399 reflections | $\Delta\rho_{\text{min}} = -0.20$ e Å ⁻³ |

Table 1

Hydrogen-bond geometry (Å, °).

Cg1 is the centroid of the C31–C36 ring.

| <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> |
|------------------------------------|-------------|---------------------|----------------------------|-------------------------------|
| C21—H21A \cdots O3 ⁱ | 0.98 | 2.57 | 3.399 (3) | 143 |
| C36—H36 \cdots O3 ⁱⁱ | 0.95 | 2.51 | 3.431 (3) | 164 |
| C22—H22B \cdots Cg1 ⁱ | 0.99 | 2.77 | 3.756 (3) | 173 |

Symmetry codes: (i) $x - \frac{1}{2}, -y + \frac{3}{2}, -z + 1$; (ii) $-x + 1, y + \frac{1}{2}, -z + \frac{3}{2}$.

Data collection: APEX2 (Bruker, 2009); cell refinement: SAINT (Bruker, 2009); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: PLATON (Spek, 2009) and Mercury (Macrae *et al.*, 2008); software used to prepare material for publication: OLEX2 (Dolomanov *et al.*, 2009).

Supporting information for this paper is available from the IUCr electronic archives (Reference: SU5049).

References

- Bruker (2009). *APEX2, SAINT and SADABS*. Bruker AXS Inc., Madison, Wisconsin, USA.
- Bugenhagen, B., Munk, A., Vill, V., Al-Jasem, Y. & Thiemann, T. (2012). *Acta Cryst.* **E68**, o2064.
- Dolomanov, O. V., Bourhis, L. J., Gildea, R. J., Howard, J. A. K. & Puschmann, H. (2009). *J. Appl. Cryst.* **42**, 339–341.
- Dong, X., Guo, J. & Wei, J. (2010). *Chin. J. Chem. Phys.* **23**, 719–725.
- Kutulya, L. A., Cherkashina, R. M., Tishchenko, V. G., Surov, Yu. N. & Polishchuk, A. G. (1983). *Zh. Obshch. Khim.* **53**, 1665–1668.
- Macrae, C. F., Bruno, I. J., Chisholm, J. A., Edgington, P. R., McCabe, P., Pidcock, E., Rodriguez-Monge, L., Taylor, R., van de Streek, J. & Wood, P. A. (2008). *J. Appl. Cryst.* **41**, 466–470.
- Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112–122.
- Spek, A. L. (2009). *Acta Cryst.* **D65**, 148–155.
- Tanaka, Y., Tsuchiya, H., Suzuki, M., Tsuda, K., Takano, J. & Kurihara, H. (1981). *Mol. Cryst. Liq. Cryst.* **68**, 113–125.
- Thiemann, T., al-Sulaibi, M., Al-Jasem, Y. & al-Hindawi, B. (2011). *Proceedings of the 15th International Electronic Conference on Synthetic Organic Chemistry*, 1–30 November 2011. Sciforum Electronic Conferences Series.
- Vora, R. A. (1976). *Curr. Sci.* **45**, 538–539.

supporting information

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Crystal structure of cholest-5-en-3 β -yl 3-(2,4-dimethoxy-3-methylphenyl)-prop-2-enoate

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S1. Structural commentary

Cholesteryl cinnamates exhibit chiral mesogenic phases. The influence of the substituents of the cinnamyl unit in these compounds on their phase transition behaviour (Vora, 1976; Kutulya *et al.*, 1983) remains of interest (Bugenhagen *et al.*, 2012). Also, their crystal packing at room temperature which can give the possibility to photodimerize the substances in the crystal (Tanaka *et al.*, 1981; Dong *et al.*, 2010) is continued to be studied. For the title compound, the authors have observed the following phase transformation sequence: Cr 162.2 Ch 229.9 I, where the numbers denote temperature of the phase transition in °C.

There are short intramolecular C—H \cdots O contacts present in the methyl (E)-3-(2,4-dimethoxy-3-methylphenyl)acrylate moiety of the title compound (Fig. 1 and Table 1).

The conformational analysis of rings A, B, C and D was carried out. It was found that rings A and C adopt a chair conformation, while ring B adopts a half-chair conformation, and ring D adopts an envelope conformation with the methyl substituted C atom as the flap.

In the crystal, molecules are arranged in separate layers parallel to (1 0 0). Within each layer, translation related molecules form columns extend along [1 0 1] with their long molecular axis collinear with this direction (Figure 3). Molecules in the neighbouring columns exhibit head to tail arrangement with C—H \cdots O interactions occurring between the 2-methoxy group (O3) of one cinnamate unit with the C6—H (H36) of the other molecule's cinnamate unit (Table 1). The neighboring layers are packed in such a manner that there are two close contact, (C21—H21A \cdots O3) and (C22—H22B \cdots π) between molecules in a head-to-tail arrangement, with a dihedral angle between the steroidal mean planes of these contacting molecules of 46.7 (2)° (Fig. 2 and Table 1). These interactions lead to the formation of a three-dimensional framework.

S2. Synthesis and crystallization

To a solution of triphenylphosphine (582 mg, 2.2 mmol) in CH₂Cl₂ (7.5 mL) is added bromotrichloromethane (900 mg, 4.5 mmol), and the resulting solution is stirred for 20 min. at rt. Thereafter, 3-(2',4'-dimethoxy-3-'methylphenyl)prop-2-enoic acid (2,4-dimethoxy-3-methylcinnamic acid, 444 mg, 2.0 mmol) is added, and the solution is heated at 323 K for 15 min. Cholest-5-en-3 β -ol (cholesterol, 386 mg, 1.0 mmol) is added, and after 20 min. Et₃N (200 mg, 2.0 mmol) is added dropwise with the help of a syringe. The reaction mixture is stirred at 318 K for 12h. Then, it is cooled, poured into water (30 mL) and extracted with CH₂Cl₂ (3 \times 15 mL). The organic phase is washed with 15 w% aq. NaOH (15 mL) and subsequently with aq. HCl (1 mL conc. HCl in 7 mL of H₂O), dried over anhydrous MgSO₄, and evaporated *in vacuo*. Column chromatography of the residue on silica gel (eluent M_tBE/hexane/CHCl₃ 1:3:1) gives the target compound (413 mg, 70%) as colorless needles; δ_{H} (400 MHz, CDCl₃) 0.68 (3H, s, CH₃), 0.86 (3H, d, 3J = 6.8 Hz, CH₃), 0.87 (3H, d, 3J = 6.8 Hz, CH₃), 0.92 (3H, d, 3J = 6.4 Hz, CH₃), 0.98 – 2.17 (26H, m), 1.05 (3H, s, CH₃), 2.41 (2H, m), 3.73 (3H, s, OCH₃),

3.85 (3H, s, OCH₃), 4.73 (1H, m), 5.41 (1H, m), 6.38 (1H, d, ³J = 16.0 Hz), 6.66 (1H, d, ³J = 8.7 Hz), 7.41 (1H, d, ³J = 8.7 Hz), 7.89 (1H, d, ³J = 16.0 Hz); δ_c (100.5 MHz, CDCl₃) 8.9, 11.9, 18.7, 19.4, 21.0, 22.6, 22.8, 23.8, 24.3, 27.9, 28.0, 28.2, 31.9, 35.8, 36.2, 36.6, 37.0, 38.3, 39.5, 39.7, 42.3, 50.0, 55.7, 56.1, 56.7, 61.5, 73.8, 106.5, 117.0, 120.2, 120.7, 122.6, 126.0, 139.8, 139.9, 158.9, 160.5, 167.0.

S3. Refinement

All H atoms were placed in calculated positions with C—H distances of 0.95 - 1.00 Å and refined as riding with $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C})$ for methyl H atoms and $= 1.2U_{\text{eq}}(\text{C})$ for other H-atoms. In the final cycles of refinement, in the absence of significant anomalous scattering effects, the Friedel pairs were merged and $\Delta f'$ set to zero.

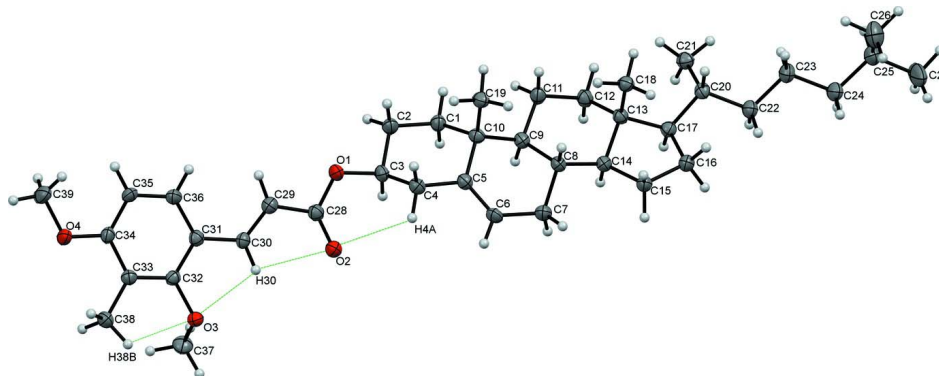


Figure 1

A view of molecular structure of the title molecule, with atom labelling. Displacement ellipsoids are shown at the 50% probability level. The short intramolecular C-H...O contacts are shown as green dashed lines (see Table 1 for details).

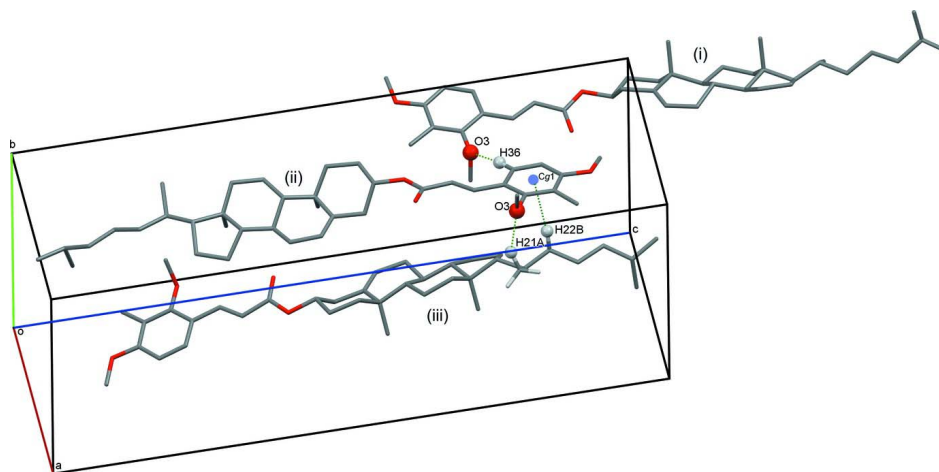
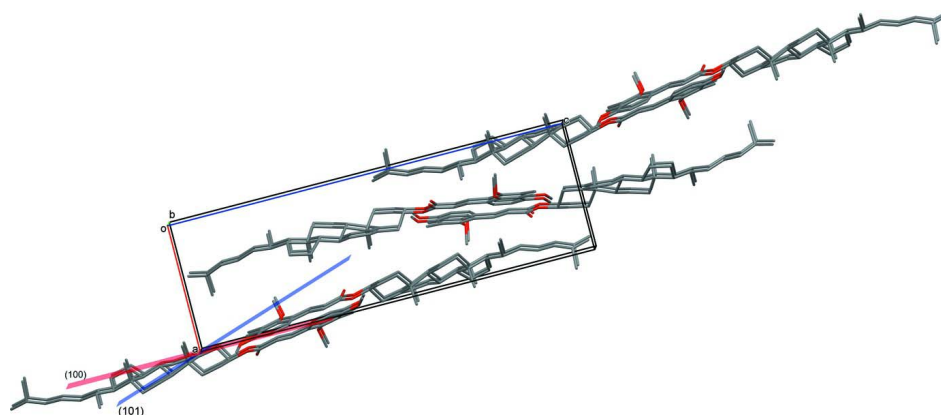


Figure 2

Intermolecular C—H...O and C—H... π (Cg1) contacts between molecules of the title compound (see Table 1 for details; symmetry codes: (i) $-x + 1, y + 1/2, -z + 3/2$; (ii) x, y, z ; (iii) $x + 1/2, -y + 3/2, -z + 1$).

**Figure 3**

A view of adjacent molecules lying in layers (three layers in this figure) parallel to (100), showing their long molecular axis which is parallel to the [101] direction.

Cholest-5-en-3 β -yl 3-(2,4-dimethoxy-3-methylphenyl)prop-2-enoate

Crystal data

$C_{39}H_{58}O_4$

$M_r = 590.85$

Orthorhombic, $P2_12_12_1$

$a = 9.4626$ (6) Å

$b = 12.2687$ (8) Å

$c = 29.6074$ (18) Å

$V = 3437.2$ (4) Å³

$Z = 4$

$F(000) = 1296$

$D_x = 1.142$ Mg m⁻³

Melting point: 503 K

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

Cell parameters from 4605 reflections

$\theta = 2.3$ – 20.4°

$\mu = 0.07$ mm⁻¹

$T = 100$ K

Needle, colourless

$0.25 \times 0.07 \times 0.07$ mm

Data collection

Bruker SMART APEX CCD area-detector diffractometer

Radiation source: micro-focus

Multi-layer monochromator

Detector resolution: 8 pixels mm⁻¹

ω and φ scans

Absorption correction: multi-scan

(*SADABS*; Bruker, 2009)

$T_{\min} = 0.702$, $T_{\max} = 0.746$

33550 measured reflections

4399 independent reflections

3563 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.052$

$\theta_{\max} = 27.5^\circ$, $\theta_{\min} = 2.2^\circ$

$h = -12 \rightarrow 12$

$k = -15 \rightarrow 15$

$l = -38 \rightarrow 38$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.046$

$wR(F^2) = 0.112$

$S = 1.03$

4399 reflections

396 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-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0454P)^2 + 1.4494P]$

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

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

$\Delta\rho_{\max} = 0.56$ e Å⁻³

$\Delta\rho_{\min} = -0.20$ e 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 (\AA^2)

| | <i>x</i> | <i>y</i> | <i>z</i> | $U_{\text{iso}}^*/U_{\text{eq}}$ |
|-----|------------|------------|--------------|----------------------------------|
| C1 | 0.3236 (3) | 0.9302 (2) | 0.48990 (8) | 0.0265 (6) |
| C10 | 0.4360 (3) | 0.8630 (2) | 0.46390 (8) | 0.0231 (6) |
| C11 | 0.3496 (3) | 0.9329 (2) | 0.38602 (8) | 0.0275 (6) |
| C12 | 0.3055 (3) | 0.9041 (2) | 0.33754 (8) | 0.0278 (6) |
| C13 | 0.4131 (3) | 0.8268 (2) | 0.31576 (8) | 0.0228 (6) |
| C14 | 0.4170 (3) | 0.7240 (2) | 0.34577 (8) | 0.0242 (6) |
| C15 | 0.5026 (3) | 0.6419 (2) | 0.31835 (8) | 0.0264 (6) |
| C16 | 0.4683 (3) | 0.6701 (2) | 0.26855 (8) | 0.0267 (6) |
| C17 | 0.3738 (3) | 0.7732 (2) | 0.26950 (8) | 0.0239 (6) |
| C18 | 0.5575 (3) | 0.8817 (2) | 0.31194 (9) | 0.0260 (6) |
| C19 | 0.5755 (3) | 0.9279 (2) | 0.45997 (9) | 0.0286 (6) |
| C2 | 0.3567 (3) | 0.9469 (2) | 0.54015 (9) | 0.0284 (6) |
| C20 | 0.3855 (3) | 0.8421 (2) | 0.22638 (8) | 0.0235 (6) |
| C21 | 0.2799 (3) | 0.9367 (2) | 0.22650 (9) | 0.0283 (6) |
| C22 | 0.3674 (3) | 0.7725 (2) | 0.18347 (8) | 0.0280 (6) |
| C23 | 0.4106 (3) | 0.8306 (2) | 0.14017 (8) | 0.0319 (7) |
| C24 | 0.4104 (4) | 0.7584 (2) | 0.09850 (8) | 0.0345 (7) |
| C25 | 0.4636 (4) | 0.8142 (3) | 0.05567 (9) | 0.0349 (7) |
| C26 | 0.6193 (4) | 0.8428 (3) | 0.05855 (11) | 0.0492 (9) |
| C27 | 0.4329 (6) | 0.7462 (4) | 0.01422 (10) | 0.0776 (16) |
| C28 | 0.4001 (3) | 0.7813 (2) | 0.64075 (9) | 0.0253 (6) |
| C29 | 0.4332 (3) | 0.8217 (2) | 0.68646 (8) | 0.0256 (6) |
| C3 | 0.3759 (3) | 0.8379 (2) | 0.56332 (8) | 0.0256 (6) |
| C30 | 0.4294 (3) | 0.7554 (2) | 0.72230 (8) | 0.0224 (5) |
| C31 | 0.4576 (3) | 0.7846 (2) | 0.76936 (8) | 0.0225 (5) |
| C32 | 0.4539 (3) | 0.7051 (2) | 0.80346 (8) | 0.0215 (5) |
| C33 | 0.4829 (3) | 0.7288 (2) | 0.84855 (8) | 0.0231 (6) |
| C34 | 0.5126 (3) | 0.8372 (2) | 0.85963 (8) | 0.0239 (6) |
| C35 | 0.5165 (3) | 0.9187 (2) | 0.82683 (8) | 0.0252 (6) |
| C36 | 0.4898 (3) | 0.8915 (2) | 0.78239 (8) | 0.0257 (6) |
| C37 | 0.2873 (3) | 0.5625 (3) | 0.80063 (10) | 0.0344 (7) |
| C38 | 0.4917 (3) | 0.6416 (2) | 0.88433 (8) | 0.0287 (6) |
| C39 | 0.5834 (4) | 0.9622 (2) | 0.91719 (10) | 0.0398 (8) |
| C4 | 0.4913 (3) | 0.7723 (2) | 0.54083 (8) | 0.0263 (6) |
| C5 | 0.4638 (3) | 0.7582 (2) | 0.49066 (8) | 0.0239 (6) |

| | | | | |
|------|------------|------------|-------------|------------|
| C6 | 0.4669 (3) | 0.6604 (2) | 0.47162 (8) | 0.0259 (6) |
| C7 | 0.4521 (3) | 0.6396 (2) | 0.42197 (8) | 0.0290 (6) |
| C8 | 0.4648 (3) | 0.7443 (2) | 0.39412 (8) | 0.0233 (6) |
| C9 | 0.3748 (3) | 0.8333 (2) | 0.41646 (8) | 0.0238 (6) |
| H11A | 0.2749 | 0.9788 | 0.3998 | 0.033* |
| H11B | 0.4373 | 0.9768 | 0.3849 | 0.033* |
| H12A | 0.2113 | 0.8690 | 0.3379 | 0.033* |
| H12B | 0.2984 | 0.9716 | 0.3194 | 0.033* |
| H14 | 0.3181 | 0.6954 | 0.3474 | 0.029* |
| H15A | 0.4737 | 0.5663 | 0.3256 | 0.032* |
| H15B | 0.6050 | 0.6499 | 0.3245 | 0.032* |
| H16A | 0.5563 | 0.6846 | 0.2515 | 0.032* |
| H16B | 0.4178 | 0.6089 | 0.2539 | 0.032* |
| H17 | 0.2736 | 0.7480 | 0.2718 | 0.029* |
| H18A | 0.5943 | 0.8968 | 0.3422 | 0.039* |
| H18B | 0.5479 | 0.9503 | 0.2952 | 0.039* |
| H18C | 0.6228 | 0.8334 | 0.2959 | 0.039* |
| H19A | 0.6409 | 0.8888 | 0.4400 | 0.043* |
| H19B | 0.6181 | 0.9356 | 0.4900 | 0.043* |
| H19C | 0.5560 | 1.0002 | 0.4474 | 0.043* |
| H1A | 0.3144 | 1.0025 | 0.4754 | 0.032* |
| H1B | 0.2312 | 0.8929 | 0.4872 | 0.032* |
| H20 | 0.4826 | 0.8742 | 0.2256 | 0.028* |
| H21A | 0.1861 | 0.9095 | 0.2349 | 0.042* |
| H21B | 0.2756 | 0.9693 | 0.1963 | 0.042* |
| H21C | 0.3103 | 0.9919 | 0.2484 | 0.042* |
| H22A | 0.4247 | 0.7054 | 0.1866 | 0.034* |
| H22B | 0.2671 | 0.7503 | 0.1809 | 0.034* |
| H23A | 0.3453 | 0.8924 | 0.1350 | 0.038* |
| H23B | 0.5066 | 0.8611 | 0.1443 | 0.038* |
| H24A | 0.3128 | 0.7323 | 0.0932 | 0.041* |
| H24B | 0.4702 | 0.6938 | 0.1045 | 0.041* |
| H25 | 0.4103 | 0.8841 | 0.0524 | 0.042* |
| H26A | 0.6355 | 0.8900 | 0.0847 | 0.074* |
| H26B | 0.6480 | 0.8809 | 0.0310 | 0.074* |
| H26C | 0.6748 | 0.7758 | 0.0618 | 0.074* |
| H27A | 0.4889 | 0.6790 | 0.0153 | 0.116* |
| H27B | 0.4578 | 0.7876 | -0.0129 | 0.116* |
| H27C | 0.3321 | 0.7278 | 0.0134 | 0.116* |
| H29 | 0.4576 | 0.8961 | 0.6905 | 0.031* |
| H2A | 0.4440 | 0.9907 | 0.5433 | 0.034* |
| H2B | 0.2784 | 0.9873 | 0.5547 | 0.034* |
| H3 | 0.2851 | 0.7962 | 0.5622 | 0.031* |
| H30 | 0.4059 | 0.6813 | 0.7166 | 0.027* |
| H35 | 0.5372 | 0.9919 | 0.8349 | 0.030* |
| H36 | 0.4935 | 0.9469 | 0.7600 | 0.031* |
| H37A | 0.2203 | 0.6075 | 0.7836 | 0.052* |
| H37B | 0.2683 | 0.5697 | 0.8330 | 0.052* |

| | | | | |
|------|------------|--------------|-------------|------------|
| H37C | 0.2767 | 0.4860 | 0.7917 | 0.052* |
| H38A | 0.5758 | 0.6538 | 0.9030 | 0.043* |
| H38B | 0.4981 | 0.5699 | 0.8699 | 0.043* |
| H38C | 0.4071 | 0.6445 | 0.9034 | 0.043* |
| H39A | 0.6663 | 0.9831 | 0.8994 | 0.060* |
| H39B | 0.6076 | 0.9631 | 0.9494 | 0.060* |
| H39C | 0.5063 | 1.0137 | 0.9115 | 0.060* |
| H4A | 0.4973 | 0.6997 | 0.5553 | 0.032* |
| H4B | 0.5830 | 0.8096 | 0.5452 | 0.032* |
| H6 | 0.4794 | 0.5991 | 0.4908 | 0.031* |
| H7A | 0.3591 | 0.6056 | 0.4160 | 0.035* |
| H7B | 0.5262 | 0.5877 | 0.4123 | 0.035* |
| H8 | 0.5659 | 0.7682 | 0.3939 | 0.028* |
| H9 | 0.2799 | 0.8001 | 0.4220 | 0.029* |
| O1 | 0.4114 (2) | 0.86254 (15) | 0.61032 (6) | 0.0278 (4) |
| O2 | 0.3647 (2) | 0.68939 (15) | 0.63125 (6) | 0.0299 (5) |
| O3 | 0.4279 (2) | 0.59773 (14) | 0.79131 (6) | 0.0237 (4) |
| O4 | 0.5397 (2) | 0.85517 (14) | 0.90460 (6) | 0.0304 (5) |

Atomic displacement parameters (Å²)

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|-----|-------------|-------------|-------------|--------------|--------------|--------------|
| C1 | 0.0330 (16) | 0.0255 (14) | 0.0212 (13) | 0.0032 (12) | 0.0000 (11) | -0.0011 (11) |
| C10 | 0.0248 (14) | 0.0223 (13) | 0.0222 (12) | 0.0015 (11) | -0.0009 (11) | 0.0022 (10) |
| C11 | 0.0340 (16) | 0.0267 (14) | 0.0219 (13) | 0.0069 (13) | 0.0003 (11) | -0.0004 (11) |
| C12 | 0.0317 (16) | 0.0291 (15) | 0.0226 (13) | 0.0047 (13) | -0.0003 (12) | 0.0012 (12) |
| C13 | 0.0242 (13) | 0.0231 (13) | 0.0211 (12) | 0.0016 (11) | 0.0006 (11) | 0.0012 (11) |
| C14 | 0.0292 (14) | 0.0220 (13) | 0.0213 (12) | -0.0011 (12) | 0.0021 (11) | 0.0002 (10) |
| C15 | 0.0348 (16) | 0.0225 (13) | 0.0218 (12) | 0.0009 (12) | 0.0019 (11) | -0.0006 (11) |
| C16 | 0.0332 (15) | 0.0240 (13) | 0.0228 (12) | 0.0006 (12) | 0.0024 (12) | -0.0017 (11) |
| C17 | 0.0245 (14) | 0.0238 (13) | 0.0234 (12) | -0.0008 (12) | 0.0003 (11) | -0.0013 (11) |
| C18 | 0.0305 (15) | 0.0230 (13) | 0.0246 (13) | -0.0021 (12) | -0.0027 (12) | 0.0028 (11) |
| C19 | 0.0339 (16) | 0.0266 (14) | 0.0254 (13) | -0.0057 (13) | -0.0012 (12) | 0.0020 (11) |
| C2 | 0.0391 (16) | 0.0229 (14) | 0.0233 (13) | 0.0057 (13) | 0.0011 (12) | 0.0006 (11) |
| C20 | 0.0217 (13) | 0.0269 (14) | 0.0219 (12) | -0.0029 (11) | -0.0013 (11) | -0.0002 (11) |
| C21 | 0.0301 (15) | 0.0274 (15) | 0.0273 (13) | -0.0013 (12) | -0.0038 (12) | 0.0016 (12) |
| C22 | 0.0324 (15) | 0.0282 (14) | 0.0233 (13) | -0.0012 (13) | -0.0023 (12) | -0.0020 (12) |
| C23 | 0.0379 (17) | 0.0343 (15) | 0.0236 (13) | -0.0024 (14) | -0.0013 (12) | -0.0033 (12) |
| C24 | 0.0469 (19) | 0.0323 (16) | 0.0243 (13) | -0.0058 (14) | -0.0044 (13) | -0.0014 (12) |
| C25 | 0.0486 (19) | 0.0333 (16) | 0.0227 (13) | -0.0015 (15) | -0.0017 (14) | 0.0005 (12) |
| C26 | 0.048 (2) | 0.066 (2) | 0.0335 (16) | 0.0039 (19) | 0.0100 (15) | -0.0044 (17) |
| C27 | 0.144 (5) | 0.067 (3) | 0.0216 (16) | -0.045 (3) | -0.004 (2) | -0.0024 (18) |
| C28 | 0.0277 (15) | 0.0250 (14) | 0.0231 (13) | 0.0035 (12) | 0.0015 (11) | 0.0016 (11) |
| C29 | 0.0283 (14) | 0.0237 (13) | 0.0249 (13) | 0.0001 (12) | 0.0012 (11) | -0.0014 (11) |
| C3 | 0.0334 (15) | 0.0259 (14) | 0.0174 (12) | -0.0022 (12) | -0.0031 (11) | 0.0014 (11) |
| C30 | 0.0232 (13) | 0.0226 (13) | 0.0215 (12) | 0.0015 (11) | 0.0028 (11) | -0.0017 (11) |
| C31 | 0.0220 (13) | 0.0233 (13) | 0.0222 (12) | 0.0018 (11) | 0.0018 (11) | -0.0003 (10) |
| C32 | 0.0223 (13) | 0.0192 (13) | 0.0231 (12) | 0.0011 (11) | 0.0013 (11) | -0.0020 (10) |

| | | | | | | |
|-----|-------------|-------------|-------------|--------------|--------------|--------------|
| C33 | 0.0253 (14) | 0.0214 (13) | 0.0227 (12) | 0.0016 (12) | 0.0003 (11) | 0.0005 (10) |
| C34 | 0.0283 (14) | 0.0240 (13) | 0.0194 (11) | 0.0028 (12) | 0.0017 (11) | -0.0032 (10) |
| C35 | 0.0344 (15) | 0.0168 (12) | 0.0245 (12) | 0.0018 (12) | -0.0009 (12) | -0.0026 (10) |
| C36 | 0.0331 (16) | 0.0211 (13) | 0.0230 (12) | 0.0024 (12) | 0.0022 (12) | 0.0018 (11) |
| C37 | 0.0279 (16) | 0.0366 (17) | 0.0387 (16) | -0.0086 (14) | 0.0036 (13) | -0.0003 (14) |
| C38 | 0.0382 (16) | 0.0252 (13) | 0.0227 (12) | -0.0004 (13) | -0.0016 (12) | 0.0026 (11) |
| C39 | 0.066 (2) | 0.0264 (15) | 0.0270 (14) | -0.0051 (16) | -0.0106 (15) | -0.0039 (12) |
| C4 | 0.0308 (15) | 0.0248 (13) | 0.0233 (12) | 0.0004 (12) | -0.0009 (12) | 0.0027 (11) |
| C5 | 0.0246 (14) | 0.0248 (13) | 0.0223 (12) | 0.0007 (12) | 0.0013 (11) | 0.0028 (11) |
| C6 | 0.0325 (15) | 0.0230 (13) | 0.0222 (12) | 0.0024 (12) | 0.0006 (11) | 0.0062 (11) |
| C7 | 0.0394 (17) | 0.0219 (14) | 0.0257 (13) | 0.0007 (13) | 0.0032 (12) | 0.0016 (11) |
| C8 | 0.0283 (14) | 0.0222 (13) | 0.0194 (11) | -0.0013 (12) | 0.0004 (11) | 0.0021 (10) |
| C9 | 0.0262 (14) | 0.0239 (13) | 0.0211 (12) | -0.0010 (12) | 0.0011 (11) | 0.0011 (11) |
| O1 | 0.0404 (12) | 0.0245 (10) | 0.0186 (8) | -0.0014 (9) | -0.0025 (8) | 0.0015 (7) |
| O2 | 0.0406 (12) | 0.0248 (10) | 0.0244 (9) | -0.0029 (9) | -0.0005 (9) | 0.0003 (8) |
| O3 | 0.0261 (10) | 0.0201 (9) | 0.0250 (9) | -0.0021 (8) | 0.0016 (8) | -0.0009 (7) |
| O4 | 0.0490 (13) | 0.0220 (9) | 0.0200 (9) | -0.0010 (10) | -0.0042 (9) | -0.0020 (7) |

Geometric parameters (Å, °)

| | | | |
|----------|-----------|----------|-----------|
| O1—C3 | 1.463 (3) | C32—O3 | 1.388 (3) |
| O1—C28 | 1.348 (3) | C10—C1 | 1.550 (4) |
| O2—C28 | 1.210 (3) | C10—C5 | 1.533 (4) |
| C3—H3 | 1.0000 | C10—C19 | 1.546 (4) |
| C3—C4 | 1.511 (4) | C17—H17 | 1.0000 |
| C3—C2 | 1.514 (4) | C1—H1A | 0.9900 |
| C30—H30 | 0.9500 | C1—H1B | 0.9900 |
| C30—C29 | 1.338 (3) | C1—C2 | 1.534 (3) |
| C30—C31 | 1.463 (3) | C5—C4 | 1.518 (3) |
| C34—C33 | 1.399 (4) | C21—H21A | 0.9800 |
| C34—C35 | 1.394 (4) | C21—H21B | 0.9800 |
| C34—O4 | 1.374 (3) | C21—H21C | 0.9800 |
| C8—H8 | 1.0000 | C12—H12A | 0.9900 |
| C8—C14 | 1.522 (3) | C12—H12B | 0.9900 |
| C8—C7 | 1.531 (3) | C12—C11 | 1.536 (4) |
| C8—C9 | 1.535 (4) | C22—H22A | 0.9900 |
| C14—H14 | 1.0000 | C22—H22B | 0.9900 |
| C14—C15 | 1.526 (4) | C22—C23 | 1.523 (4) |
| C14—C13 | 1.544 (4) | C4—H4A | 0.9900 |
| C28—C29 | 1.475 (4) | C4—H4B | 0.9900 |
| C29—H29 | 0.9500 | C2—H2A | 0.9900 |
| C15—H15A | 0.9900 | C2—H2B | 0.9900 |
| C15—H15B | 0.9900 | C24—H24A | 0.9900 |
| C15—C16 | 1.549 (3) | C24—H24B | 0.9900 |
| C7—H7A | 0.9900 | C24—C23 | 1.519 (4) |
| C7—H7B | 0.9900 | C11—H11A | 0.9900 |
| C7—C6 | 1.499 (3) | C11—H11B | 0.9900 |
| C6—H6 | 0.9500 | C23—H23A | 0.9900 |

| | | | |
|-------------|-----------|---------------|-----------|
| C6—C5 | 1.326 (4) | C23—H23B | 0.9900 |
| C31—C36 | 1.401 (4) | C26—H26A | 0.9800 |
| C31—C32 | 1.404 (3) | C26—H26B | 0.9800 |
| C33—C32 | 1.394 (3) | C26—H26C | 0.9800 |
| C33—C38 | 1.508 (4) | O4—C39 | 1.426 (3) |
| C20—H20 | 1.0000 | O3—C37 | 1.426 (3) |
| C20—C17 | 1.535 (3) | C18—H18A | 0.9800 |
| C20—C21 | 1.532 (4) | C18—H18B | 0.9800 |
| C20—C22 | 1.540 (3) | C18—H18C | 0.9800 |
| C13—C17 | 1.564 (3) | C19—H19A | 0.9800 |
| C13—C12 | 1.534 (4) | C19—H19B | 0.9800 |
| C13—C18 | 1.527 (4) | C19—H19C | 0.9800 |
| C35—H35 | 0.9500 | C38—H38A | 0.9800 |
| C35—C36 | 1.381 (3) | C38—H38B | 0.9800 |
| C16—H16A | 0.9900 | C38—H38C | 0.9800 |
| C16—H16B | 0.9900 | C37—H37A | 0.9800 |
| C16—C17 | 1.549 (4) | C37—H37B | 0.9800 |
| C36—H36 | 0.9500 | C37—H37C | 0.9800 |
| C25—H25 | 1.0000 | C39—H39A | 0.9800 |
| C25—C24 | 1.526 (4) | C39—H39B | 0.9800 |
| C25—C26 | 1.516 (5) | C39—H39C | 0.9800 |
| C25—C27 | 1.512 (4) | C27—H27A | 0.9800 |
| C9—H9 | 1.0000 | C27—H27B | 0.9800 |
| C9—C10 | 1.562 (3) | C27—H27C | 0.9800 |
| C9—C11 | 1.537 (3) | | |
| | | | |
| C28—O1—C3 | 117.7 (2) | C16—C17—H17 | 107.2 |
| O1—C3—H3 | 109.5 | C10—C1—H1A | 108.7 |
| O1—C3—C4 | 111.3 (2) | C10—C1—H1B | 108.7 |
| O1—C3—C2 | 106.0 (2) | H1A—C1—H1B | 107.6 |
| C4—C3—H3 | 109.5 | C2—C1—C10 | 114.4 (2) |
| C4—C3—C2 | 110.9 (2) | C2—C1—H1A | 108.7 |
| C2—C3—H3 | 109.5 | C2—C1—H1B | 108.7 |
| C29—C30—H30 | 116.5 | C6—C5—C10 | 122.9 (2) |
| C29—C30—C31 | 127.0 (2) | C6—C5—C4 | 121.0 (2) |
| C31—C30—H30 | 116.5 | C4—C5—C10 | 116.1 (2) |
| C35—C34—C33 | 121.6 (2) | C20—C21—H21A | 109.5 |
| O4—C34—C33 | 114.7 (2) | C20—C21—H21B | 109.5 |
| O4—C34—C35 | 123.7 (2) | C20—C21—H21C | 109.5 |
| C14—C8—H8 | 109.0 | H21A—C21—H21B | 109.5 |
| C14—C8—C7 | 110.2 (2) | H21A—C21—H21C | 109.5 |
| C14—C8—C9 | 110.9 (2) | H21B—C21—H21C | 109.5 |
| C7—C8—H8 | 109.0 | C13—C12—H12A | 109.5 |
| C7—C8—C9 | 108.8 (2) | C13—C12—H12B | 109.5 |
| C9—C8—H8 | 109.0 | C13—C12—C11 | 110.8 (2) |
| C8—C14—H14 | 106.9 | H12A—C12—H12B | 108.1 |
| C8—C14—C15 | 116.8 (2) | C11—C12—H12A | 109.5 |
| C8—C14—C13 | 114.5 (2) | C11—C12—H12B | 109.5 |

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|---------------|-----------|---------------|-----------|
| C15—C14—H14 | 106.9 | C20—C22—H22A | 108.8 |
| C15—C14—C13 | 104.2 (2) | C20—C22—H22B | 108.8 |
| C13—C14—H14 | 106.9 | H22A—C22—H22B | 107.7 |
| O1—C28—C29 | 110.4 (2) | C23—C22—C20 | 113.9 (2) |
| O2—C28—O1 | 123.8 (2) | C23—C22—H22A | 108.8 |
| O2—C28—C29 | 125.8 (2) | C23—C22—H22B | 108.8 |
| C30—C29—C28 | 121.2 (2) | C3—C4—C5 | 111.6 (2) |
| C30—C29—H29 | 119.4 | C3—C4—H4A | 109.3 |
| C28—C29—H29 | 119.4 | C3—C4—H4B | 109.3 |
| C14—C15—H15A | 110.9 | C5—C4—H4A | 109.3 |
| C14—C15—H15B | 110.9 | C5—C4—H4B | 109.3 |
| C14—C15—C16 | 104.3 (2) | H4A—C4—H4B | 108.0 |
| H15A—C15—H15B | 108.9 | C3—C2—C1 | 110.3 (2) |
| C16—C15—H15A | 110.9 | C3—C2—H2A | 109.6 |
| C16—C15—H15B | 110.9 | C3—C2—H2B | 109.6 |
| C8—C7—H7A | 109.2 | C1—C2—H2A | 109.6 |
| C8—C7—H7B | 109.2 | C1—C2—H2B | 109.6 |
| H7A—C7—H7B | 107.9 | H2A—C2—H2B | 108.1 |
| C6—C7—C8 | 112.2 (2) | C25—C24—H24A | 108.7 |
| C6—C7—H7A | 109.2 | C25—C24—H24B | 108.7 |
| C6—C7—H7B | 109.2 | H24A—C24—H24B | 107.6 |
| C7—C6—H6 | 117.7 | C23—C24—C25 | 114.4 (2) |
| C5—C6—C7 | 124.7 (2) | C23—C24—H24A | 108.7 |
| C5—C6—H6 | 117.7 | C23—C24—H24B | 108.7 |
| C36—C31—C30 | 122.1 (2) | C9—C11—H11A | 108.8 |
| C36—C31—C32 | 117.2 (2) | C9—C11—H11B | 108.8 |
| C32—C31—C30 | 120.7 (2) | C12—C11—C9 | 114.0 (2) |
| C34—C33—C38 | 119.9 (2) | C12—C11—H11A | 108.8 |
| C32—C33—C34 | 117.5 (2) | C12—C11—H11B | 108.8 |
| C32—C33—C38 | 122.4 (2) | H11A—C11—H11B | 107.6 |
| C17—C20—H20 | 107.6 | C22—C23—H23A | 108.7 |
| C17—C20—C22 | 111.9 (2) | C22—C23—H23B | 108.7 |
| C21—C20—H20 | 107.6 | C24—C23—C22 | 114.2 (2) |
| C21—C20—C17 | 111.6 (2) | C24—C23—H23A | 108.7 |
| C21—C20—C22 | 110.5 (2) | C24—C23—H23B | 108.7 |
| C22—C20—H20 | 107.6 | H23A—C23—H23B | 107.6 |
| C14—C13—C17 | 99.5 (2) | C25—C26—H26A | 109.5 |
| C12—C13—C14 | 106.2 (2) | C25—C26—H26B | 109.5 |
| C12—C13—C17 | 118.0 (2) | C25—C26—H26C | 109.5 |
| C18—C13—C14 | 112.5 (2) | H26A—C26—H26B | 109.5 |
| C18—C13—C17 | 109.5 (2) | H26A—C26—H26C | 109.5 |
| C18—C13—C12 | 110.6 (2) | H26B—C26—H26C | 109.5 |
| C34—C35—H35 | 120.5 | C34—O4—C39 | 117.1 (2) |
| C36—C35—C34 | 119.1 (2) | C32—O3—C37 | 113.8 (2) |
| C36—C35—H35 | 120.5 | C13—C18—H18A | 109.5 |
| C15—C16—H16A | 110.4 | C13—C18—H18B | 109.5 |
| C15—C16—H16B | 110.4 | C13—C18—H18C | 109.5 |
| H16A—C16—H16B | 108.6 | H18A—C18—H18B | 109.5 |

| | | | |
|-------------------------|-----------|-------------------------|------------|
| C17—C16—C15 | 106.7 (2) | H18A—C18—H18C | 109.5 |
| C17—C16—H16A | 110.4 | H18B—C18—H18C | 109.5 |
| C17—C16—H16B | 110.4 | C10—C19—H19A | 109.5 |
| C31—C36—H36 | 119.1 | C10—C19—H19B | 109.5 |
| C35—C36—C31 | 121.9 (2) | C10—C19—H19C | 109.5 |
| C35—C36—H36 | 119.1 | H19A—C19—H19B | 109.5 |
| C24—C25—H25 | 107.3 | H19A—C19—H19C | 109.5 |
| C26—C25—H25 | 107.3 | H19B—C19—H19C | 109.5 |
| C26—C25—C24 | 112.2 (3) | C33—C38—H38A | 109.5 |
| C27—C25—H25 | 107.3 | C33—C38—H38B | 109.5 |
| C27—C25—C24 | 111.3 (3) | C33—C38—H38C | 109.5 |
| C27—C25—C26 | 111.1 (3) | H38A—C38—H38B | 109.5 |
| C8—C9—H9 | 106.3 | H38A—C38—H38C | 109.5 |
| C8—C9—C10 | 110.4 (2) | H38B—C38—H38C | 109.5 |
| C8—C9—C11 | 113.6 (2) | O3—C37—H37A | 109.5 |
| C10—C9—H9 | 106.3 | O3—C37—H37B | 109.5 |
| C11—C9—H9 | 106.3 | O3—C37—H37C | 109.5 |
| C11—C9—C10 | 113.5 (2) | H37A—C37—H37B | 109.5 |
| C33—C32—C31 | 122.7 (2) | H37A—C37—H37C | 109.5 |
| O3—C32—C31 | 118.5 (2) | H37B—C37—H37C | 109.5 |
| O3—C32—C33 | 118.7 (2) | O4—C39—H39A | 109.5 |
| C1—C10—C9 | 108.5 (2) | O4—C39—H39B | 109.5 |
| C5—C10—C9 | 109.4 (2) | O4—C39—H39C | 109.5 |
| C5—C10—C1 | 107.9 (2) | H39A—C39—H39B | 109.5 |
| C5—C10—C19 | 108.9 (2) | H39A—C39—H39C | 109.5 |
| C19—C10—C9 | 111.6 (2) | H39B—C39—H39C | 109.5 |
| C19—C10—C1 | 110.5 (2) | C25—C27—H27A | 109.5 |
| C20—C17—C13 | 118.6 (2) | C25—C27—H27B | 109.5 |
| C20—C17—C16 | 113.1 (2) | C25—C27—H27C | 109.5 |
| C20—C17—H17 | 107.2 | H27A—C27—H27B | 109.5 |
| C13—C17—H17 | 107.2 | H27A—C27—H27C | 109.5 |
| C16—C17—C13 | 102.8 (2) | H27B—C27—H27C | 109.5 |
| | | | |
| C(1)—C(2)—C(3)—O(1) | 178.2 (2) | C(5)—C(10)—C(19)—H(19A) | 69 |
| C(1)—C(2)—C(3)—C(4) | 57.2 (3) | C(5)—C(10)—C(19)—H(19B) | -51 |
| C(1)—C(2)—C(3)—H(3) | -64 | C(5)—C(10)—C(19)—H(19C) | -171 |
| C(1)—C(10)—C(19)—H(19A) | -172 | C(6)—C(5)—C(10)—C(1) | 132.6 (3) |
| C(1)—C(10)—C(19)—H(19B) | 68 | C(6)—C(5)—C(10)—C(9) | 14.8 (4) |
| C(1)—C(10)—C(19)—H(19C) | -52 | C(6)—C(5)—C(10)—C(19) | -107.5 (3) |
| C(10)—C(1)—C(2)—C(3) | -56.7 (3) | C(6)—C(7)—C(8)—C(9) | -45.6 (3) |
| C(10)—C(5)—C(6)—C(7) | 3.6 (5) | C(6)—C(7)—C(8)—C(14) | -167.4 (2) |
| C(10)—C(9)—C(11)—C(12) | 173.4 (2) | C(6)—C(7)—C(8)—H(8) | 73 |
| C(10)—C(1)—C(2)—H(2A) | 64 | C(7)—C(8)—C(9)—C(10) | 65.3 (3) |
| C(10)—C(1)—C(2)—H(2B) | -177 | C(7)—C(8)—C(9)—C(11) | -166.0 (2) |
| C(10)—C(5)—C(6)—H(6) | -176 | C(7)—C(8)—C(14)—C(13) | 174.9 (2) |
| C(10)—C(9)—C(11)—H(11A) | -65 | C(7)—C(8)—C(14)—C(15) | -62.9 (3) |
| C(10)—C(9)—C(11)—H(11B) | 52 | C(7)—C(8)—C(9)—H(9) | -50 |
| C(11)—C(9)—C(10)—C(1) | 65.2 (3) | C(7)—C(8)—C(14)—H(14) | 57 |

| | | | |
|--------------------------|------------|---------------------------|------------|
| C(11)—C(9)—C(10)—C(5) | -177.3 (2) | C(8)—C(9)—C(10)—C(1) | -166.0 (2) |
| C(11)—C(9)—C(10)—C(19) | -56.7 (3) | C(8)—C(9)—C(10)—C(5) | -48.6 (3) |
| C(11)—C(12)—C(13)—C(14) | 59.5 (3) | C(8)—C(9)—C(10)—C(19) | 72.1 (3) |
| C(11)—C(12)—C(13)—C(17) | 170.1 (2) | C(8)—C(9)—C(11)—C(12) | 46.3 (3) |
| C(11)—C(12)—C(13)—C(18) | -62.8 (3) | C(8)—C(14)—C(15)—C(16) | -159.4 (2) |
| C(12)—C(13)—C(14)—C(8) | -61.6 (3) | C(8)—C(9)—C(11)—H(11A) | 168 |
| C(12)—C(13)—C(14)—C(15) | 169.6 (2) | C(8)—C(9)—C(11)—H(11B) | -75 |
| C(12)—C(13)—C(17)—C(16) | -156.9 (2) | C(8)—C(14)—C(15)—H(15A) | 81 |
| C(12)—C(13)—C(17)—C(20) | 77.4 (3) | C(8)—C(14)—C(15)—H(15B) | -40 |
| C(12)—C(13)—C(14)—H(14) | 57 | C(9)—C(8)—C(14)—C(13) | 54.4 (3) |
| C(12)—C(13)—C(17)—H(17) | -44 | C(9)—C(8)—C(14)—C(15) | 176.6 (2) |
| C(12)—C(13)—C(18)—H(18A) | 64 | C(9)—C(11)—C(12)—C(13) | -54.8 (3) |
| C(12)—C(13)—C(18)—H(18B) | -56 | C(9)—C(8)—C(14)—H(14) | -64 |
| C(12)—C(13)—C(18)—H(18C) | -176 | C(9)—C(10)—C(19)—H(19A) | -52 |
| C(13)—C(14)—C(15)—C(16) | -32.0 (3) | C(9)—C(10)—C(19)—H(19B) | -172 |
| C(13)—C(17)—C(20)—C(21) | -65.2 (3) | C(9)—C(10)—C(19)—H(19C) | 68 |
| C(13)—C(17)—C(20)—C(22) | 170.5 (2) | C(9)—C(11)—C(12)—H(12A) | 66 |
| C(13)—C(14)—C(15)—H(15A) | -151 | C(9)—C(11)—C(12)—H(12B) | -176 |
| C(13)—C(14)—C(15)—H(15B) | 88 | H(11A)—C(11)—C(12)—C(13) | -176 |
| C(13)—C(17)—C(20)—H(20) | 53 | H(11A)—C(11)—C(12)—H(12A) | -55 |
| C(14)—C(8)—C(9)—C(10) | -173.3 (2) | H(11A)—C(11)—C(12)—H(12B) | 63 |
| C(14)—C(8)—C(9)—C(11) | -44.6 (3) | H(11B)—C(11)—C(12)—C(13) | 67 |
| C(14)—C(13)—C(17)—C(16) | -42.6 (2) | H(11B)—C(11)—C(12)—H(12A) | -172 |
| C(14)—C(13)—C(17)—C(20) | -168.3 (2) | H(11B)—C(11)—C(12)—H(12B) | -54 |
| C(14)—C(15)—C(16)—C(17) | 4.4 (3) | H(12A)—C(12)—C(13)—C(14) | -61 |
| C(14)—C(8)—C(9)—H(9) | 72 | H(12A)—C(12)—C(13)—C(17) | 49 |
| C(14)—C(13)—C(17)—H(17) | 70 | H(12A)—C(12)—C(13)—C(18) | 176 |
| C(14)—C(13)—C(18)—H(18A) | -54 | H(12B)—C(12)—C(13)—C(14) | -180 |
| C(14)—C(13)—C(18)—H(18B) | -174 | H(12B)—C(12)—C(13)—C(17) | -69 |
| C(14)—C(13)—C(18)—H(18C) | 66 | H(12B)—C(12)—C(13)—C(18) | 58 |
| C(14)—C(15)—C(16)—H(16A) | 124 | H(14)—C(14)—C(15)—C(16) | 81 |
| C(14)—C(15)—C(16)—H(16B) | -116 | H(14)—C(14)—C(15)—H(15A) | -38 |
| C(15)—C(16)—C(17)—C(13) | 24.1 (3) | H(14)—C(14)—C(15)—H(15B) | -159 |
| C(15)—C(16)—C(17)—C(20) | 153.3 (2) | H(15A)—C(15)—C(16)—C(17) | 124 |
| C(15)—C(16)—C(17)—H(17) | -89 | H(15A)—C(15)—C(16)—H(16A) | -116 |
| C(16)—C(17)—C(20)—C(21) | 174.2 (2) | H(15A)—C(15)—C(16)—H(16B) | 4 |
| C(16)—C(17)—C(20)—C(22) | 49.9 (3) | H(15B)—C(15)—C(16)—C(17) | -115 |
| C(16)—C(17)—C(20)—H(20) | -68 | H(15B)—C(15)—C(16)—H(16A) | 5 |
| C(17)—C(13)—C(14)—C(8) | 175.3 (2) | H(15B)—C(15)—C(16)—H(16B) | 125 |
| C(17)—C(13)—C(14)—C(15) | 46.5 (3) | H(16A)—C(16)—C(17)—C(13) | -96 |
| C(17)—C(20)—C(22)—C(23) | -166.9 (2) | H(16A)—C(16)—C(17)—C(20) | 33 |
| C(17)—C(13)—C(14)—H(14) | -66 | H(16A)—C(16)—C(17)—H(17) | 151 |
| C(17)—C(13)—C(18)—H(18A) | -164 | H(16B)—C(16)—C(17)—C(13) | 144 |
| C(17)—C(13)—C(18)—H(18B) | 76 | H(16B)—C(16)—C(17)—C(20) | -87 |
| C(17)—C(13)—C(18)—H(18C) | -44 | H(16B)—C(16)—C(17)—H(17) | 31 |
| C(17)—C(20)—C(21)—H(21A) | -48 | H(17)—C(17)—C(20)—C(21) | 56 |
| C(17)—C(20)—C(21)—H(21B) | -168 | H(17)—C(17)—C(20)—C(22) | -68 |
| C(17)—C(20)—C(21)—H(21C) | 72 | H(17)—C(17)—C(20)—H(20) | 174 |

| | | | |
|--------------------------|------------|---------------------------|------|
| C(17)—C(20)—C(22)—H(22A) | -45 | H(1A)—C(1)—C(2)—C(3) | -178 |
| C(17)—C(20)—C(22)—H(22B) | 72 | H(1A)—C(1)—C(2)—H(2A) | -58 |
| C(18)—C(13)—C(14)—C(8) | 59.5 (3) | H(1A)—C(1)—C(2)—H(2B) | 61 |
| C(18)—C(13)—C(14)—C(15) | -69.3 (3) | H(1A)—C(1)—C(10)—C(5) | 172 |
| C(18)—C(13)—C(17)—C(16) | 75.4 (2) | H(1A)—C(1)—C(10)—C(9) | -69 |
| C(18)—C(13)—C(17)—C(20) | -50.3 (3) | H(1A)—C(1)—C(10)—C(19) | 53 |
| C(18)—C(13)—C(14)—H(14) | 178 | H(1B)—C(1)—C(2)—C(3) | 65 |
| C(18)—C(13)—C(17)—H(17) | -172 | H(1B)—C(1)—C(2)—H(2A) | -174 |
| C(2)—C(1)—C(10)—C(5) | 50.6 (3) | H(1B)—C(1)—C(2)—H(2B) | -56 |
| C(2)—C(1)—C(10)—C(9) | 169.0 (2) | H(1B)—C(1)—C(10)—C(5) | -71 |
| C(2)—C(1)—C(10)—C(19) | -68.4 (3) | H(1B)—C(1)—C(10)—C(9) | 47 |
| C(2)—C(3)—C(4)—C(5) | -55.3 (3) | H(1B)—C(1)—C(10)—C(19) | 170 |
| C(2)—C(3)—C(4)—H(4A) | -176 | H(20)—C(20)—C(21)—H(21A) | -166 |
| C(2)—C(3)—C(4)—H(4B) | 66 | H(20)—C(20)—C(21)—H(21B) | 74 |
| C(20)—C(22)—C(23)—C(24) | 172.8 (3) | H(20)—C(20)—C(21)—H(21C) | -46 |
| C(20)—C(22)—C(23)—H(23A) | -66 | H(20)—C(20)—C(22)—C(23) | -49 |
| C(20)—C(22)—C(23)—H(23B) | 51 | H(20)—C(20)—C(22)—H(22A) | 73 |
| C(21)—C(20)—C(22)—C(23) | 68.1 (3) | H(20)—C(20)—C(22)—H(22B) | -170 |
| C(21)—C(20)—C(22)—H(22A) | -170 | H(22A)—C(22)—C(23)—C(24) | 51 |
| C(21)—C(20)—C(22)—H(22B) | -53 | H(22A)—C(22)—C(23)—H(23A) | 173 |
| C(22)—C(23)—C(24)—C(25) | -175.9 (3) | H(22A)—C(22)—C(23)—H(23B) | -70 |
| C(22)—C(20)—C(21)—H(21A) | 77 | H(22B)—C(22)—C(23)—C(24) | -66 |
| C(22)—C(20)—C(21)—H(21B) | -43 | H(22B)—C(22)—C(23)—H(23A) | 56 |
| C(22)—C(20)—C(21)—H(21C) | -163 | H(22B)—C(22)—C(23)—H(23B) | 173 |
| C(22)—C(23)—C(24)—H(24A) | 62 | H(23A)—C(23)—C(24)—C(25) | 63 |
| C(22)—C(23)—C(24)—H(24B) | -54 | H(23A)—C(23)—C(24)—H(24A) | -59 |
| C(23)—C(24)—C(25)—C(26) | 66.4 (4) | H(23A)—C(23)—C(24)—H(24B) | -176 |
| C(23)—C(24)—C(25)—C(27) | -168.4 (3) | H(23B)—C(23)—C(24)—C(25) | -54 |
| C(23)—C(24)—C(25)—H(25) | -51 | H(23B)—C(23)—C(24)—H(24A) | -176 |
| C(24)—C(25)—C(26)—H(26A) | -58 | H(23B)—C(23)—C(24)—H(24B) | 67 |
| C(24)—C(25)—C(26)—H(26B) | -178 | H(24A)—C(24)—C(25)—C(26) | -172 |
| C(24)—C(25)—C(26)—H(26C) | 62 | H(24A)—C(24)—C(25)—C(27) | -47 |
| C(24)—C(25)—C(27)—H(27A) | -67 | H(24A)—C(24)—C(25)—H(25) | 70 |
| C(24)—C(25)—C(27)—H(27B) | 173 | H(24B)—C(24)—C(25)—C(26) | -55 |
| C(24)—C(25)—C(27)—H(27C) | 53 | H(24B)—C(24)—C(25)—C(27) | 70 |
| C(26)—C(25)—C(27)—H(27A) | 59 | H(24B)—C(24)—C(25)—H(25) | -173 |
| C(26)—C(25)—C(27)—H(27B) | -61 | H(25)—C(25)—C(26)—H(26A) | 60 |
| C(26)—C(25)—C(27)—H(27C) | 179 | H(25)—C(25)—C(26)—H(26B) | -60 |
| C(27)—C(25)—C(26)—H(26A) | 177 | H(25)—C(25)—C(26)—H(26C) | 180 |
| C(27)—C(25)—C(26)—H(26B) | 57 | H(25)—C(25)—C(27)—H(27A) | 176 |
| C(27)—C(25)—C(26)—H(26C) | -63 | H(25)—C(25)—C(27)—H(27B) | 56 |
| C(28)—O(1)—C(3)—C(2) | 164.1 (2) | H(25)—C(25)—C(27)—H(27C) | -64 |
| C(28)—O(1)—C(3)—C(4) | -75.2 (3) | H(29)—C(29)—C(30)—C(31) | 1 |
| C(28)—C(29)—C(30)—C(31) | -179.1 (3) | H(29)—C(29)—C(30)—H(30) | -179 |
| C(28)—O(1)—C(3)—H(3) | 46 | H(2A)—C(2)—C(3)—O(1) | 57 |
| C(28)—C(29)—C(30)—H(30) | 1 | H(2A)—C(2)—C(3)—C(4) | -63 |
| C(29)—C(30)—C(31)—C(32) | -178.0 (3) | H(2A)—C(2)—C(3)—H(3) | 176 |
| C(29)—C(30)—C(31)—C(36) | 1.7 (5) | H(2B)—C(2)—C(3)—O(1) | -61 |

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|--------------------------|------------|-------------------------|------------|
| C(3)—O(1)—C(28)—O(2) | 1.3 (4) | H(2B)—C(2)—C(3)—C(4) | 178 |
| C(3)—O(1)—C(28)—C(29) | -177.3 (2) | H(2B)—C(2)—C(3)—H(3) | 57 |
| C(3)—C(4)—C(5)—C(6) | -128.5 (3) | H(3)—C(3)—C(4)—C(5) | 66 |
| C(3)—C(4)—C(5)—C(10) | 52.8 (3) | H(3)—C(3)—C(4)—H(4A) | -55 |
| C(30)—C(31)—C(32)—O(3) | 2.3 (4) | H(3)—C(3)—C(4)—H(4B) | -173 |
| C(30)—C(31)—C(32)—C(33) | 178.7 (3) | H(30)—C(30)—C(31)—C(32) | 2 |
| C(30)—C(31)—C(36)—C(35) | -179.9 (3) | H(30)—C(30)—C(31)—C(36) | -178 |
| C(30)—C(31)—C(36)—H(36) | 0 | H(35)—C(35)—C(36)—C(31) | -179 |
| C(31)—C(32)—C(33)—C(34) | 1.9 (4) | H(35)—C(35)—C(36)—H(36) | 1 |
| C(31)—C(32)—C(33)—C(38) | -174.4 (3) | H(4A)—C(4)—C(5)—C(6) | -7 |
| C(32)—C(31)—C(36)—C(35) | -0.2 (4) | H(4A)—C(4)—C(5)—C(10) | 174 |
| C(32)—C(33)—C(34)—O(4) | 179.8 (2) | H(4B)—C(4)—C(5)—C(6) | 111 |
| C(32)—C(33)—C(34)—C(35) | -1.5 (4) | H(4B)—C(4)—C(5)—C(10) | -68 |
| C(32)—O(3)—C(37)—H(37A) | 61 | H(6)—C(6)—C(7)—C(8) | -168 |
| C(32)—O(3)—C(37)—H(37B) | -59 | H(6)—C(6)—C(7)—H(7A) | 71 |
| C(32)—O(3)—C(37)—H(37C) | -179 | H(6)—C(6)—C(7)—H(7B) | -47 |
| C(32)—C(31)—C(36)—H(36) | 180 | H(7A)—C(7)—C(8)—C(9) | 76 |
| C(32)—C(33)—C(38)—H(38A) | 134 | H(7A)—C(7)—C(8)—C(14) | -46 |
| C(32)—C(33)—C(38)—H(38B) | 14 | H(7A)—C(7)—C(8)—H(8) | -166 |
| C(32)—C(33)—C(38)—H(38C) | -106 | H(7B)—C(7)—C(8)—C(9) | -167 |
| C(33)—C(34)—C(35)—C(36) | 0.3 (4) | H(7B)—C(7)—C(8)—C(14) | 72 |
| C(33)—C(34)—C(35)—H(35) | -180 | H(7B)—C(7)—C(8)—H(8) | -48 |
| C(34)—C(35)—C(36)—C(31) | 0.6 (4) | H(8)—C(8)—C(9)—C(10) | -53 |
| C(34)—O(4)—C(39)—H(39A) | -55 | H(8)—C(8)—C(9)—C(11) | 75 |
| C(34)—O(4)—C(39)—H(39B) | -175 | H(8)—C(8)—C(9)—H(9) | -168 |
| C(34)—O(4)—C(39)—H(39C) | 65 | H(8)—C(8)—C(14)—C(13) | -66 |
| C(34)—C(33)—C(38)—H(38A) | -42 | H(8)—C(8)—C(14)—C(15) | 57 |
| C(34)—C(33)—C(38)—H(38B) | -162 | H(8)—C(8)—C(14)—H(14) | 176 |
| C(34)—C(33)—C(38)—H(38C) | 78 | H(9)—C(9)—C(10)—C(1) | -51 |
| C(34)—C(35)—C(36)—H(36) | -179 | H(9)—C(9)—C(10)—C(5) | 66 |
| C(36)—C(31)—C(32)—O(3) | -177.4 (2) | H(9)—C(9)—C(10)—C(19) | -173 |
| C(36)—C(31)—C(32)—C(33) | -1.1 (4) | H(9)—C(9)—C(11)—C(12) | -70 |
| C(37)—O(3)—C(32)—C(31) | -103.0 (3) | H(9)—C(9)—C(11)—H(11A) | 51 |
| C(37)—O(3)—C(32)—C(33) | 80.5 (3) | H(9)—C(9)—C(11)—H(11B) | 168 |
| C(38)—C(33)—C(34)—O(4) | -3.9 (4) | O(1)—C(3)—C(4)—C(5) | -173.1 (2) |
| C(38)—C(33)—C(34)—C(35) | 174.9 (3) | O(1)—C(28)—C(29)—C(30) | -179.8 (3) |
| C(39)—O(4)—C(34)—C(33) | 174.4 (3) | O(1)—C(3)—C(4)—H(4A) | 66 |
| C(39)—O(4)—C(34)—C(35) | -4.4 (4) | O(1)—C(3)—C(4)—H(4B) | -52 |
| C(4)—C(5)—C(6)—C(7) | -175.1 (3) | O(1)—C(28)—C(29)—H(29) | 0 |
| C(4)—C(5)—C(10)—C(1) | -48.7 (3) | O(2)—C(28)—C(29)—C(30) | 1.7 (5) |
| C(4)—C(5)—C(10)—C(9) | -166.5 (2) | O(2)—C(28)—C(29)—H(29) | -178 |
| C(4)—C(5)—C(10)—C(19) | 71.2 (3) | O(3)—C(32)—C(33)—C(34) | 178.3 (2) |
| C(4)—C(5)—C(6)—H(6) | 5 | O(3)—C(32)—C(33)—C(38) | 2.0 (4) |
| C(5)—C(6)—C(7)—C(8) | 12.4 (4) | O(4)—C(34)—C(35)—C(36) | 178.9 (3) |
| C(5)—C(6)—C(7)—H(7A) | -109 | O(4)—C(34)—C(35)—H(35) | -1 |
| C(5)—C(6)—C(7)—H(7B) | 133 | | |

Hydrogen-bond geometry (Å, °)

Cg1 is the centroid of the C31–C36 ring.

| <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> |
|--|-------------|---------------------|----------------------------|-------------------------------|
| C4—H4 <i>A</i> \cdots O2 | 0.99 | 2.58 | 3.104 (3) | 113 |
| C30—H30 \cdots O2 | 0.95 | 2.56 | 2.881 (3) | 100 |
| C30—H30 \cdots O3 | 0.95 | 2.45 | 2.814 (3) | 103 |
| C38—H38 <i>B</i> \cdots O3 | 0.98 | 2.44 | 2.870 (3) | 106 |
| C21—H21 <i>A</i> \cdots O3 ⁱ | 0.98 | 2.57 | 3.399 (3) | 143 |
| C36—H36 \cdots O3 ⁱⁱ | 0.95 | 2.51 | 3.431 (3) | 164 |
| C22—H22 <i>B</i> \cdots Cg1 ⁱ | 0.99 | 2.77 | 3.756 (3) | 173 |

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