

5,22-Stigmastadien-3 β -yl *p*-toluenesulfonate

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Key indicators: single-crystal X-ray study; $T = 100 \text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.003 \text{ \AA}$;
 R factor = 0.035; wR factor = 0.089; data-to-parameter ratio = 17.8.

The asymmetric unit of the title compound [systematic name: (3*S*,8*S*,9*S*,10*R*,13*R*,14*S*,17*R*)-17-[(*E*,2*R*,5*S*)-5-ethyl-6-methylhept-3-en-2-yl]-10,13-dimethyl-2,3,4,7,8,9,11,12,14,15,16,17-dodecahydro-1*H*-cyclopenta[*a*]phenanthren-3-yl *p*-toluenesulfonate], $C_{36}H_{54}O_3S$, comprises two independent molecules that differ significantly in terms of the relative orientations of the peripheral groups; the conformation about the $\text{C}=\text{C}$ bond of the side chain is *E*. In the crystal, molecules associate into linear supramolecular chains aligned along the *a* axis via $\text{C}-\text{H}\cdots\text{O}$ interactions.

Related literature

For the use of 5,22-stigmastadien-3 β -yl *p*-toluenesulfonate, see: Partridge *et al.* (1974); Khripach *et al.* (2002); Foley *et al.* (2010); Ketuly *et al.* (1997). For the synthesis, see: Foley *et al.* (2010).

Data collection

Bruker SMART APEX CCD diffractometer
Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996)
 (SADABS) ; Sheldrick, 1996)
 $T_{\min} = 0.669$, $T_{\max} = 0.746$

15647 measured reflections
13115 independent reflections
12382 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.018$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.035$
 $wR(F^2) = 0.089$
 $S = 1.02$
13115 reflections
735 parameters
3 restraints

H-atom parameters constrained
 $\Delta\rho_{\max} = 0.35 \text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.37 \text{ e \AA}^{-3}$
Absolute structure: Flack (1983),
5643 Friedel pairs
Flack parameter: 0.02 (4)

Table 1
Hydrogen-bond geometry (\AA , $^\circ$).

| $D-\text{H}\cdots A$ | $D-\text{H}$ | $\text{H}\cdots A$ | $D\cdots A$ | $D-\text{H}\cdots A$ |
|-----------------------------------|--------------|--------------------|-------------|----------------------|
| C7—H7 \cdots O3 ⁱ | 0.95 | 2.49 | 3.217 (2) | 134 |
| C13—H13B \cdots O6 | 0.99 | 2.54 | 3.519 (2) | 172 |
| C40—H40 \cdots O5 ⁱⁱ | 0.95 | 2.48 | 3.193 (2) | 131 |
| C42—H42A \cdots O2 | 0.99 | 2.56 | 3.548 (2) | 175 |
| C44A—H44C \cdots O2 | 0.98 | 2.54 | 3.390 (2) | 145 |

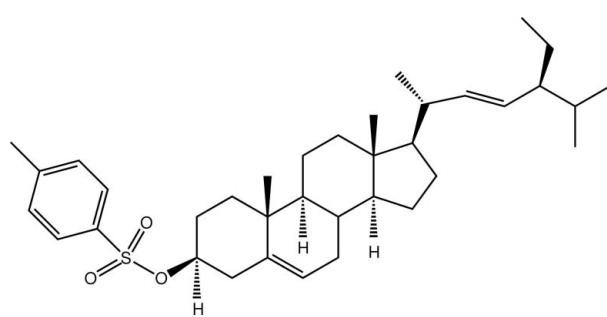
Symmetry codes: (i) $x - 1, y, z$; (ii) $x + 1, y, z$.

Data collection: *APEX2* (Bruker, 2008); cell refinement: *SAINT* (Bruker, 2008); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3* (Farrugia, 1997), *DIAMOND* (Brandenburg, 2006) and *Qmol* (Gans & Shalloway, 2001); software used to prepare material for publication: *publCIF* (Westrip, 2010).

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB5438).

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Experimental

Crystal data

$C_{36}H_{54}O_3S$
 $M_r = 566.86$
Triclinic, $P\bar{1}$
 $a = 7.0361(1) \text{ \AA}$
 $b = 11.2350(1) \text{ \AA}$
 $c = 21.1550(2) \text{ \AA}$

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supporting information

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S1. Comment

The title compound, (I), a stigmasterol tosylate, has been used as a precursor for the synthesis of cholesterol and other sterols with variable side-chains (Partridge *et al.* 1974; Khrapach *et al.* 2002; Foley *et al.* 2010) and for the determination of absolute configuration (Ketuly *et al.* 1997). Herein, (I) has been characterised by crystallography and shown to crystallise with two independent molecules in the crystallographic asymmetric unit, Figs 1 and 2. While the central steroid residues in the independent molecules are virtually superimposable, as seen from Fig. 3, the peripheral groups have quite different orientations. These differences are quantified in the values of the C5–S1–O1–C8 and C38–S2–O4–C41 torsion angles of 67.92 (13) and -63.53 (14) $^{\circ}$, respectively, which indicate that the tosylate groups lie to either side of the respective molecule. The differences in the orientation of the terminal *iso*-propyl group are best quantified in the C27–C28–C29–C30, C31 torsion angles of 45.2 (2) and 168.45 (16) $^{\circ}$, respectively, compared with 58.2 (2) and -66.8 (2) $^{\circ}$ for C60–C61–C62–C63,C64, respectively. The conformation about the C=C [C26=C27 = 1.322 (3) Å and C59=C60 = 1.324 (2) Å] double bond of the side-chain in each molecule is *E*.

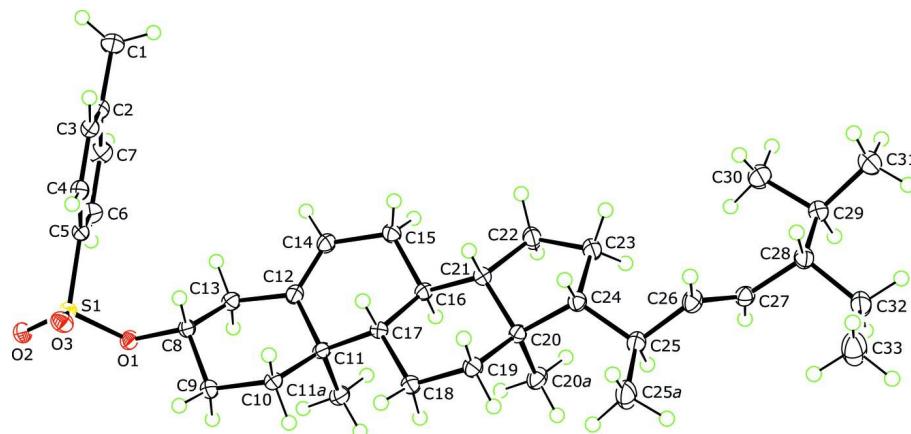
In the crystal structure, the molecules associate into linear supramolecular chains aligned along the *a* axis mediated by C–H \cdots O interactions, Fig. 2 and Table 1. The chains inter-digitate along the *b* axis as shown in Fig. 5.

S2. Experimental

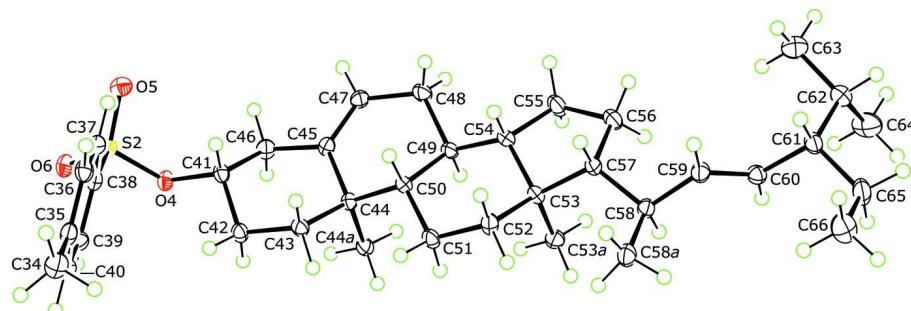
Stigmasterol (40 g, 97 mmol) was dissolved in dried and redistilled pyridine (250 ml) and *p*-toluenesulfonyl chloride (30 g, 157.4 mmol) added, and the solution was then stirred at room temperature for 28 h. The reaction mixture was mixed with 5% aqueous sodium bicarbonate and the solid precipitate filtered, washed with water, dried at room temperature, and recovered (52 g). The crude product was recrystallized from acetone, yielding fine crystals of stigmasterol tosylate (45.8 g), m.pt. 407–410 K. The compound was further recrystallised and colourless blocks of (I) were grown from n-hexane:carbon tetrachloride (1/1), m.pt. 415–417 K [Lit. 415–417 K (Foley *et al.*, 2010)].

S3. Refinement

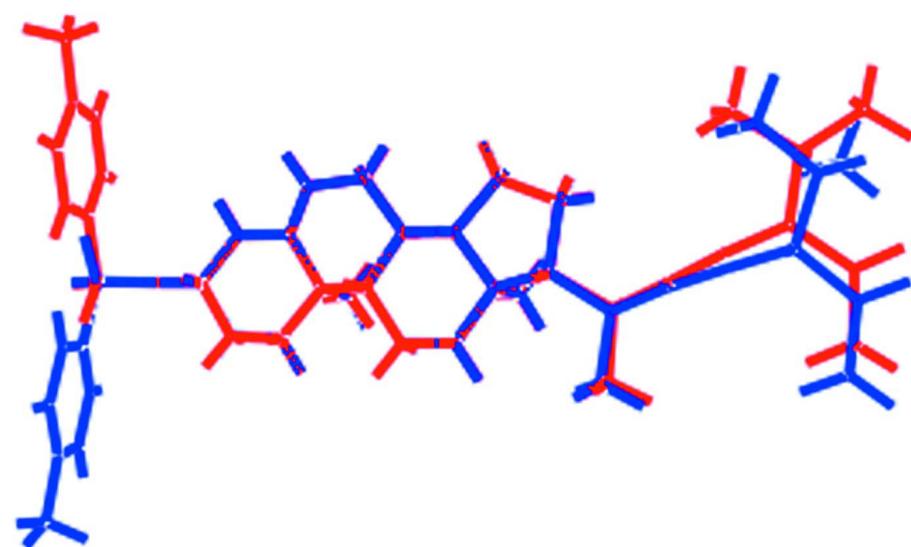
The H atoms were geometrically placed (C–H = 0.95–1.00 Å) and refined as riding with $U_{iso}(\text{H}) = 1.2\text{--}1.5U_{eq}(\text{C})$.

**Figure 1**

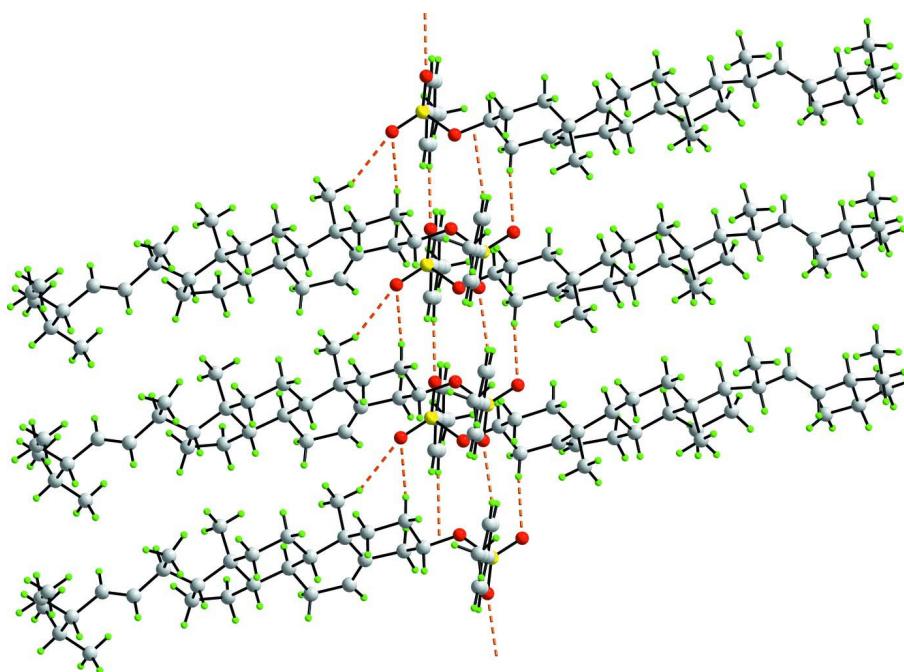
The molecular structure of the first independent molecule in (I) displacement ellipsoids at the 50% probability level.

**Figure 2**

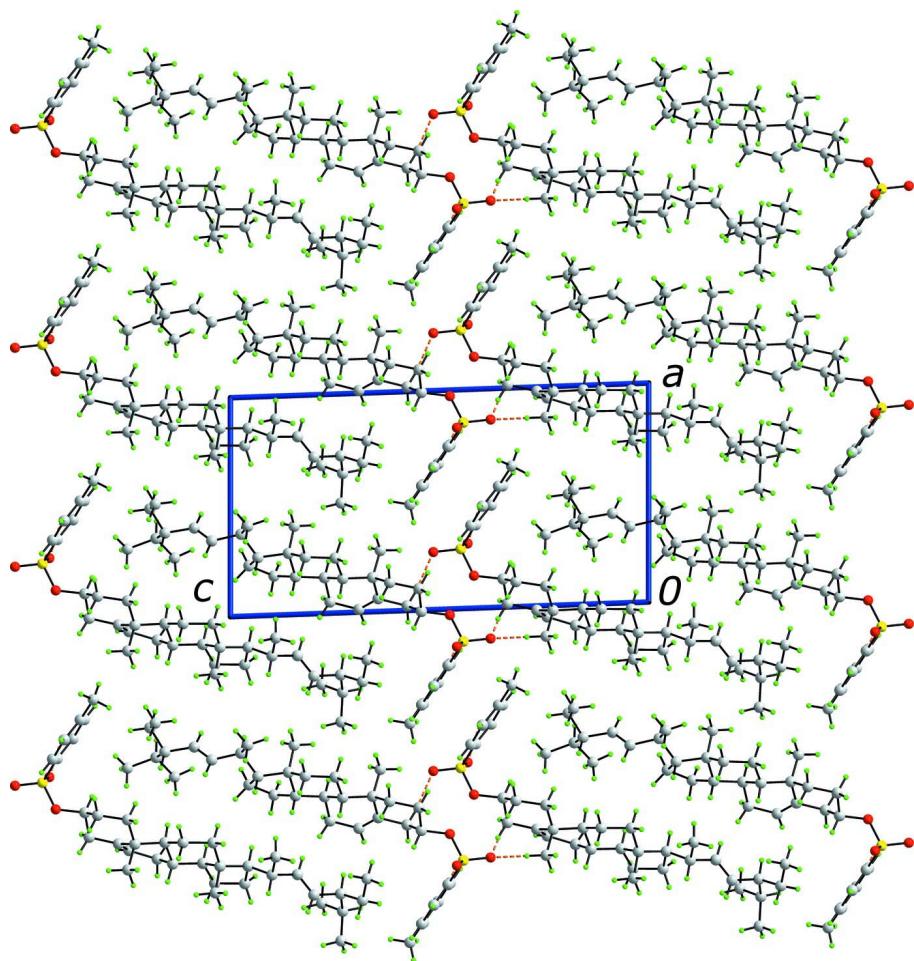
The molecular structure of the second independent molecule in (I) showing displacement ellipsoids at the 50% probability level.

**Figure 3**

Overlap diagram highlighting the different conformations of the first (red) and second (blue) independent molecules in (I).

**Figure 4**

A view of a supramolecular chain in (I) aligned along the a axis. The C–H \cdots O contacts are shown as orange dashed lines. Colour code: O, red; C, grey; and H, green.

**Figure 5**

A view in projection down the a axis of the unit cell contents in (I) showing the inter-digitation of supramolecular chains. The C–H···O contacts are shown as orange dashed lines, respectively. Colour code: O, red; C, grey; and H, green.

(3*S*,8*S*,9*S*,10*R*,13*R*,14*S*,17*R*)-17-[*(E*,2*R*,5*S*)-5-ethyl-6-methylhept-3-en-2-yl]-10,13-dimethyl-2,3,4,7,8,9,11,12,14,15,16,17-dodecahydro-1*H*-cyclopenta[*a*]phenanthren-3-yl *p*-toluenesulfonate

Crystal data

$C_{36}H_{54}O_3S$
 $M_r = 566.86$
Triclinic, $P\bar{1}$
Hall symbol: P 1
 $a = 7.0361 (1)$ Å
 $b = 11.2350 (1)$ Å
 $c = 21.1550 (2)$ Å
 $\alpha = 90.777 (1)^\circ$
 $\beta = 96.166 (1)^\circ$
 $\gamma = 101.153 (1)^\circ$
 $V = 1630.23 (3)$ Å³

$Z = 2$
 $F(000) = 620$
 $D_x = 1.155 \text{ Mg m}^{-3}$
Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
Cell parameters from 8182 reflections
 $\theta = 2.6\text{--}30.5^\circ$
 $\mu = 0.13 \text{ mm}^{-1}$
 $T = 100 \text{ K}$
Block, colourless
 $0.49 \times 0.37 \times 0.27 \text{ mm}$

Data collection

Bruker SMART APEX CCD
diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
 ω scans
Absorption correction: multi-scan
(*SADABS*; Sheldrick, 1996)
 $T_{\min} = 0.669$, $T_{\max} = 0.746$

15647 measured reflections
13115 independent reflections
12382 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.018$
 $\theta_{\max} = 27.5^\circ$, $\theta_{\min} = 1.9^\circ$
 $h = -9 \rightarrow 9$
 $k = -14 \rightarrow 14$
 $l = -27 \rightarrow 27$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.035$
 $wR(F^2) = 0.089$
 $S = 1.02$
13115 reflections
735 parameters
3 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
 $w = 1/[\sigma^2(F_o^2) + (0.049P)^2 + 0.2642P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.35 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.37 \text{ e } \text{\AA}^{-3}$
Absolute structure: Flack (1983), 5643 Friedel
pairs
Absolute structure parameter: 0.02 (4)

Special details

Geometry. All s.u.'s (except the s.u. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell s.u.'s are taken into account individually in the estimation of s.u.'s in distances, angles and torsion angles; correlations between s.u.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell s.u.'s is used for estimating s.u.'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 $F^2 > 2\sigma(F^2)$ 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> | $U_{\text{iso}}^*/U_{\text{eq}}$ |
|-----|--------------|--------------|--------------|----------------------------------|
| S1 | 0.69443 (5) | 0.84972 (3) | 0.44298 (2) | 0.01550 (9) |
| O1 | 0.67495 (19) | 0.97242 (11) | 0.47537 (6) | 0.0185 (3) |
| O2 | 0.62261 (19) | 0.85861 (12) | 0.37793 (6) | 0.0202 (3) |
| O3 | 0.88690 (18) | 0.82684 (12) | 0.45847 (6) | 0.0213 (3) |
| C1 | 0.1261 (3) | 0.48353 (19) | 0.57090 (10) | 0.0283 (5) |
| H1A | 0.1484 | 0.5009 | 0.6169 | 0.042* |
| H1B | 0.1450 | 0.4011 | 0.5619 | 0.042* |
| H1C | -0.0074 | 0.4903 | 0.5551 | 0.042* |
| C2 | 0.2679 (3) | 0.57305 (17) | 0.53824 (9) | 0.0208 (4) |
| C3 | 0.4643 (3) | 0.56629 (17) | 0.54289 (9) | 0.0190 (4) |
| H3 | 0.5088 | 0.5034 | 0.5663 | 0.023* |
| C4 | 0.5962 (3) | 0.65025 (16) | 0.51380 (8) | 0.0174 (4) |
| H4 | 0.7301 | 0.6452 | 0.5172 | 0.021* |
| C5 | 0.5300 (3) | 0.74157 (16) | 0.47974 (8) | 0.0155 (3) |
| C6 | 0.3344 (3) | 0.74989 (18) | 0.47399 (10) | 0.0214 (4) |

| | | | | |
|------|------------|--------------|--------------|------------|
| H6 | 0.2903 | 0.8125 | 0.4503 | 0.026* |
| C7 | 0.2052 (3) | 0.66572 (18) | 0.50327 (10) | 0.0240 (4) |
| H7 | 0.0713 | 0.6709 | 0.4996 | 0.029* |
| C8 | 0.7474 (3) | 0.99334 (17) | 0.54366 (8) | 0.0172 (4) |
| H8 | 0.7888 | 0.9189 | 0.5611 | 0.021* |
| C9 | 0.9196 (3) | 1.09800 (17) | 0.54931 (9) | 0.0206 (4) |
| H9A | 0.8814 | 1.1693 | 0.5282 | 0.025* |
| H9B | 1.0263 | 1.0754 | 0.5278 | 0.025* |
| C10 | 0.9908 (3) | 1.13062 (17) | 0.61942 (9) | 0.0193 (4) |
| H10A | 1.0969 | 1.2032 | 0.6220 | 0.023* |
| H10B | 1.0467 | 1.0630 | 0.6382 | 0.023* |
| C11 | 0.8321 (3) | 1.15659 (15) | 0.66011 (8) | 0.0171 (4) |
| C11A | 0.7752 (3) | 1.27858 (16) | 0.64174 (9) | 0.0223 (4) |
| H11A | 0.7340 | 1.2768 | 0.5959 | 0.034* |
| H11B | 0.8878 | 1.3448 | 0.6524 | 0.034* |
| H11C | 0.6678 | 1.2917 | 0.6652 | 0.034* |
| C12 | 0.6484 (3) | 1.05841 (16) | 0.64686 (9) | 0.0172 (4) |
| C13 | 0.5805 (3) | 1.02109 (17) | 0.57757 (9) | 0.0198 (4) |
| H13A | 0.4748 | 0.9483 | 0.5752 | 0.024* |
| H13B | 0.5277 | 1.0874 | 0.5561 | 0.024* |
| C14 | 0.5425 (3) | 1.01280 (16) | 0.69279 (8) | 0.0192 (3) |
| H14 | 0.4307 | 0.9513 | 0.6808 | 0.023* |
| C15 | 0.5868 (3) | 1.05133 (16) | 0.76192 (8) | 0.0190 (3) |
| H15A | 0.6230 | 0.9829 | 0.7863 | 0.023* |
| H15B | 0.4675 | 1.0697 | 0.7775 | 0.023* |
| C16 | 0.7520 (3) | 1.16293 (15) | 0.77506 (8) | 0.0162 (3) |
| H16 | 0.6978 | 1.2374 | 0.7652 | 0.019* |
| C17 | 0.9115 (3) | 1.15600 (16) | 0.73162 (8) | 0.0161 (3) |
| H17 | 0.9478 | 1.0749 | 0.7383 | 0.019* |
| C18 | 1.0995 (3) | 1.25072 (17) | 0.74971 (9) | 0.0205 (4) |
| H18A | 1.0768 | 1.3311 | 0.7359 | 0.025* |
| H18B | 1.2035 | 1.2301 | 0.7261 | 0.025* |
| C19 | 1.1718 (3) | 1.26021 (17) | 0.82118 (9) | 0.0204 (4) |
| H19A | 1.2166 | 1.1848 | 0.8339 | 0.024* |
| H19B | 1.2843 | 1.3287 | 0.8295 | 0.024* |
| C20 | 1.0109 (3) | 1.27989 (15) | 0.86124 (8) | 0.0162 (3) |
| C20A | 0.9540 (3) | 1.40312 (16) | 0.84761 (9) | 0.0226 (4) |
| H20A | 1.0693 | 1.4680 | 0.8565 | 0.034* |
| H20B | 0.8548 | 1.4158 | 0.8748 | 0.034* |
| H20C | 0.9016 | 1.4041 | 0.8028 | 0.034* |
| C21 | 0.8384 (3) | 1.17209 (16) | 0.84454 (8) | 0.0167 (3) |
| H21 | 0.8924 | 1.0969 | 0.8523 | 0.020* |
| C22 | 0.7035 (3) | 1.17900 (18) | 0.89584 (9) | 0.0234 (4) |
| H22A | 0.6270 | 1.0977 | 0.9032 | 0.028* |
| H22B | 0.6127 | 1.2342 | 0.8838 | 0.028* |
| C23 | 0.8437 (3) | 1.22925 (17) | 0.95564 (9) | 0.0217 (4) |
| H23A | 0.8421 | 1.1665 | 0.9881 | 0.026* |
| H23B | 0.8038 | 1.3007 | 0.9742 | 0.026* |

| | | | | |
|------|--------------|--------------|--------------|-------------|
| C24 | 1.0514 (3) | 1.26601 (17) | 0.93432 (9) | 0.0172 (4) |
| H24 | 1.1180 | 1.1955 | 0.9412 | 0.021* |
| C25 | 1.1738 (3) | 1.37487 (16) | 0.97497 (8) | 0.0210 (4) |
| H25 | 1.1100 | 1.4463 | 0.9670 | 0.025* |
| C25A | 1.3849 (3) | 1.4103 (2) | 0.95873 (10) | 0.0347 (5) |
| H25A | 1.4560 | 1.4774 | 0.9873 | 0.052* |
| H25B | 1.3844 | 1.4357 | 0.9146 | 0.052* |
| H25C | 1.4488 | 1.3403 | 0.9640 | 0.052* |
| C26 | 1.1761 (3) | 1.34890 (17) | 1.04477 (9) | 0.0207 (4) |
| H26 | 1.2346 | 1.2838 | 1.0599 | 0.025* |
| C27 | 1.1020 (3) | 1.41072 (16) | 1.08631 (9) | 0.0188 (4) |
| H27 | 1.0390 | 1.4731 | 1.0699 | 0.023* |
| C28 | 1.1069 (3) | 1.39278 (16) | 1.15671 (8) | 0.0198 (4) |
| H28 | 1.1668 | 1.3206 | 1.1670 | 0.024* |
| C29 | 0.9004 (3) | 1.36906 (17) | 1.17799 (8) | 0.0216 (4) |
| H29 | 0.8534 | 1.4476 | 1.1767 | 0.026* |
| C30 | 0.7557 (3) | 1.2771 (2) | 1.13373 (10) | 0.0328 (5) |
| H30A | 0.8060 | 1.2021 | 1.1302 | 0.049* |
| H30B | 0.7384 | 1.3109 | 1.0915 | 0.049* |
| H30C | 0.6300 | 1.2594 | 1.1511 | 0.049* |
| C31 | 0.8994 (3) | 1.3254 (2) | 1.24636 (10) | 0.0323 (5) |
| H31A | 0.7700 | 1.3225 | 1.2601 | 0.048* |
| H31B | 0.9967 | 1.3818 | 1.2746 | 0.048* |
| H31C | 0.9309 | 1.2442 | 1.2481 | 0.048* |
| C32 | 1.2370 (3) | 1.50536 (19) | 1.19232 (9) | 0.0277 (4) |
| H32A | 1.1836 | 1.5780 | 1.1799 | 0.033* |
| H32B | 1.2307 | 1.4964 | 1.2386 | 0.033* |
| C33 | 1.4489 (3) | 1.5262 (3) | 1.17963 (12) | 0.0451 (6) |
| H33A | 1.5061 | 1.4574 | 1.1946 | 0.068* |
| H33B | 1.5207 | 1.6009 | 1.2022 | 0.068* |
| H33C | 1.4568 | 1.5338 | 1.1339 | 0.068* |
| S2 | 0.30849 (5) | 1.27220 (3) | 0.44943 (2) | 0.01603 (9) |
| O4 | 0.32421 (19) | 1.14814 (11) | 0.41769 (6) | 0.0179 (3) |
| O5 | 0.11769 (19) | 1.29670 (12) | 0.43282 (6) | 0.0219 (3) |
| O6 | 0.3765 (2) | 1.26274 (12) | 0.51495 (6) | 0.0215 (3) |
| C34 | 0.8898 (3) | 1.64290 (19) | 0.32767 (10) | 0.0264 (4) |
| H34A | 1.0067 | 1.6706 | 0.3577 | 0.040* |
| H34B | 0.8316 | 1.7125 | 0.3147 | 0.040* |
| H34C | 0.9255 | 1.6041 | 0.2901 | 0.040* |
| C35 | 0.7448 (3) | 1.55291 (17) | 0.35912 (9) | 0.0203 (4) |
| C36 | 0.5459 (3) | 1.55371 (17) | 0.34939 (9) | 0.0198 (4) |
| H36 | 0.5015 | 1.6139 | 0.3238 | 0.024* |
| C37 | 0.4116 (3) | 1.46780 (17) | 0.37652 (9) | 0.0182 (4) |
| H37 | 0.2764 | 1.4690 | 0.3695 | 0.022* |
| C38 | 0.4769 (3) | 1.38034 (16) | 0.41391 (9) | 0.0164 (4) |
| C39 | 0.6743 (3) | 1.37796 (17) | 0.42510 (9) | 0.0192 (4) |
| H39 | 0.7182 | 1.3183 | 0.4512 | 0.023* |
| C40 | 0.8067 (3) | 1.46471 (18) | 0.39740 (9) | 0.0213 (4) |

| | | | | |
|------|-------------|--------------|--------------|------------|
| H40 | 0.9419 | 1.4637 | 0.4048 | 0.026* |
| C41 | 0.2679 (3) | 1.12867 (16) | 0.34818 (8) | 0.0170 (4) |
| H41 | 0.2172 | 1.2001 | 0.3306 | 0.020* |
| C42 | 0.4461 (3) | 1.11455 (17) | 0.31752 (9) | 0.0179 (4) |
| H42A | 0.5007 | 1.0467 | 0.3366 | 0.021* |
| H42B | 0.5467 | 1.1897 | 0.3250 | 0.021* |
| C43 | 0.3909 (3) | 1.08909 (16) | 0.24600 (9) | 0.0179 (4) |
| H43A | 0.5085 | 1.0780 | 0.2266 | 0.022* |
| H43B | 0.3480 | 1.1610 | 0.2271 | 0.022* |
| C44 | 0.2286 (2) | 0.97669 (16) | 0.22859 (8) | 0.0154 (3) |
| C44A | 0.3072 (3) | 0.86021 (16) | 0.24517 (9) | 0.0183 (4) |
| H44A | 0.4214 | 0.8581 | 0.2227 | 0.027* |
| H44B | 0.2057 | 0.7888 | 0.2321 | 0.027* |
| H44C | 0.3443 | 0.8600 | 0.2912 | 0.027* |
| C45 | 0.0589 (3) | 0.98035 (16) | 0.26753 (8) | 0.0160 (3) |
| C46 | 0.1093 (3) | 1.01536 (17) | 0.33770 (9) | 0.0193 (4) |
| H46A | 0.1538 | 0.9475 | 0.3604 | 0.023* |
| H46B | -0.0088 | 1.0300 | 0.3556 | 0.023* |
| C47 | -0.1263 (2) | 0.94840 (16) | 0.24258 (8) | 0.0195 (4) |
| H47 | -0.2239 | 0.9535 | 0.2697 | 0.023* |
| C48 | -0.1933 (3) | 0.90488 (18) | 0.17491 (9) | 0.0224 (4) |
| H48A | -0.2561 | 0.9667 | 0.1527 | 0.027* |
| H48B | -0.2928 | 0.8293 | 0.1748 | 0.027* |
| C49 | -0.0279 (2) | 0.88027 (17) | 0.13799 (8) | 0.0169 (4) |
| H49 | 0.0021 | 0.7995 | 0.1499 | 0.020* |
| C50 | 0.1550 (2) | 0.97888 (16) | 0.15671 (8) | 0.0152 (3) |
| H50 | 0.1138 | 1.0586 | 0.1502 | 0.018* |
| C51 | 0.3161 (3) | 0.97594 (17) | 0.11341 (8) | 0.0195 (4) |
| H51A | 0.3752 | 0.9043 | 0.1234 | 0.023* |
| H51B | 0.4192 | 1.0493 | 0.1231 | 0.023* |
| C52 | 0.2445 (3) | 0.97042 (16) | 0.04174 (8) | 0.0192 (4) |
| H52A | 0.2008 | 1.0467 | 0.0301 | 0.023* |
| H52B | 0.3534 | 0.9627 | 0.0170 | 0.023* |
| C53 | 0.0754 (2) | 0.86221 (15) | 0.02491 (8) | 0.0154 (3) |
| C53A | 0.1438 (3) | 0.74110 (16) | 0.03621 (9) | 0.0198 (4) |
| H53A | 0.1851 | 0.7351 | 0.0816 | 0.030* |
| H53B | 0.2536 | 0.7378 | 0.0118 | 0.030* |
| H53C | 0.0363 | 0.6734 | 0.0224 | 0.030* |
| C54 | -0.0868 (2) | 0.87808 (16) | 0.06625 (8) | 0.0175 (4) |
| H54 | -0.1178 | 0.9596 | 0.0566 | 0.021* |
| C55 | -0.2647 (3) | 0.78459 (19) | 0.03754 (9) | 0.0265 (4) |
| H55A | -0.3873 | 0.8112 | 0.0444 | 0.032* |
| H55B | -0.2656 | 0.7044 | 0.0565 | 0.032* |
| C56 | -0.2399 (3) | 0.77885 (19) | -0.03430 (9) | 0.0254 (4) |
| H56A | -0.3447 | 0.8107 | -0.0595 | 0.030* |
| H56B | -0.2456 | 0.6940 | -0.0489 | 0.030* |
| C57 | -0.0379 (3) | 0.85800 (17) | -0.04245 (9) | 0.0179 (4) |
| H57 | -0.0587 | 0.9421 | -0.0512 | 0.021* |

| | | | | |
|------|-------------|--------------|---------------|------------|
| C58 | 0.0537 (3) | 0.81453 (16) | -0.09941 (8) | 0.0186 (3) |
| H58 | 0.0752 | 0.7303 | -0.0915 | 0.022* |
| C58A | 0.2503 (3) | 0.89545 (18) | -0.10828 (9) | 0.0248 (4) |
| H58A | 0.2919 | 0.8736 | -0.1489 | 0.037* |
| H58B | 0.3478 | 0.8838 | -0.0734 | 0.037* |
| H58C | 0.2370 | 0.9807 | -0.1082 | 0.037* |
| C59 | -0.0795 (3) | 0.81142 (16) | -0.16047 (8) | 0.0195 (4) |
| H59 | -0.1195 | 0.8850 | -0.1721 | 0.023* |
| C60 | -0.1459 (3) | 0.71560 (17) | -0.19935 (9) | 0.0189 (4) |
| H60 | -0.1182 | 0.6399 | -0.1857 | 0.023* |
| C61 | -0.2616 (3) | 0.71673 (16) | -0.26345 (9) | 0.0207 (4) |
| H61 | -0.2784 | 0.8022 | -0.2703 | 0.025* |
| C62 | -0.4672 (3) | 0.63669 (18) | -0.26825 (10) | 0.0252 (4) |
| H62 | -0.5354 | 0.6497 | -0.3108 | 0.030* |
| C63 | -0.5848 (3) | 0.6760 (2) | -0.21817 (11) | 0.0367 (5) |
| H63A | -0.7194 | 0.6306 | -0.2254 | 0.055* |
| H63B | -0.5839 | 0.7631 | -0.2211 | 0.055* |
| H63C | -0.5268 | 0.6595 | -0.1758 | 0.055* |
| C64 | -0.4669 (3) | 0.50155 (19) | -0.26452 (12) | 0.0360 (5) |
| H64A | -0.4080 | 0.4846 | -0.2223 | 0.054* |
| H64B | -0.3912 | 0.4774 | -0.2971 | 0.054* |
| H64C | -0.6011 | 0.4556 | -0.2718 | 0.054* |
| C65 | -0.1447 (3) | 0.68280 (19) | -0.31588 (9) | 0.0284 (4) |
| H65A | -0.0975 | 0.6076 | -0.3044 | 0.034* |
| H65B | -0.2323 | 0.6655 | -0.3562 | 0.034* |
| C66 | 0.0285 (3) | 0.7812 (2) | -0.32658 (10) | 0.0336 (5) |
| H66A | -0.0170 | 0.8557 | -0.3387 | 0.050* |
| H66B | 0.0962 | 0.7544 | -0.3607 | 0.050* |
| H66C | 0.1183 | 0.7971 | -0.2873 | 0.050* |

Atomic displacement parameters (\AA^2)

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|-----|-------------|-------------|-------------|--------------|--------------|--------------|
| S1 | 0.0165 (2) | 0.0165 (2) | 0.0130 (2) | 0.00202 (16) | 0.00175 (15) | 0.00057 (16) |
| O1 | 0.0251 (7) | 0.0152 (6) | 0.0144 (7) | 0.0033 (5) | -0.0008 (5) | 0.0006 (5) |
| O2 | 0.0240 (7) | 0.0214 (7) | 0.0139 (7) | 0.0021 (5) | 0.0004 (5) | 0.0000 (5) |
| O3 | 0.0172 (7) | 0.0258 (7) | 0.0209 (7) | 0.0030 (5) | 0.0039 (5) | 0.0028 (6) |
| C1 | 0.0279 (11) | 0.0257 (10) | 0.0296 (11) | -0.0027 (8) | 0.0108 (9) | 0.0005 (9) |
| C2 | 0.0235 (10) | 0.0180 (9) | 0.0194 (10) | -0.0013 (7) | 0.0056 (8) | -0.0040 (7) |
| C3 | 0.0246 (10) | 0.0154 (9) | 0.0165 (9) | 0.0023 (7) | 0.0026 (7) | -0.0002 (7) |
| C4 | 0.0179 (9) | 0.0180 (9) | 0.0160 (9) | 0.0037 (7) | 0.0009 (7) | -0.0007 (7) |
| C5 | 0.0164 (9) | 0.0157 (8) | 0.0138 (9) | 0.0008 (7) | 0.0027 (7) | -0.0001 (7) |
| C6 | 0.0155 (9) | 0.0222 (10) | 0.0267 (10) | 0.0047 (7) | 0.0007 (7) | 0.0030 (8) |
| C7 | 0.0142 (9) | 0.0255 (10) | 0.0320 (11) | 0.0027 (7) | 0.0033 (8) | 0.0000 (8) |
| C8 | 0.0215 (9) | 0.0165 (9) | 0.0120 (9) | 0.0006 (7) | -0.0001 (7) | 0.0002 (7) |
| C9 | 0.0237 (10) | 0.0206 (9) | 0.0149 (9) | -0.0023 (7) | 0.0029 (7) | -0.0007 (7) |
| C10 | 0.0208 (9) | 0.0180 (9) | 0.0166 (9) | -0.0026 (7) | 0.0031 (7) | -0.0013 (7) |
| C11 | 0.0198 (9) | 0.0130 (8) | 0.0161 (9) | -0.0013 (7) | 0.0001 (7) | 0.0012 (6) |

| | | | | | | |
|------|-------------|-------------|-------------|--------------|--------------|---------------|
| C11A | 0.0281 (10) | 0.0172 (9) | 0.0202 (9) | 0.0015 (7) | 0.0013 (7) | 0.0008 (7) |
| C12 | 0.0188 (9) | 0.0156 (8) | 0.0167 (9) | 0.0033 (7) | 0.0005 (7) | -0.0010 (7) |
| C13 | 0.0212 (9) | 0.0178 (9) | 0.0182 (9) | -0.0008 (7) | 0.0008 (7) | 0.0000 (7) |
| C14 | 0.0173 (8) | 0.0190 (8) | 0.0194 (9) | -0.0007 (6) | 0.0008 (7) | -0.0024 (7) |
| C15 | 0.0176 (9) | 0.0210 (8) | 0.0174 (9) | 0.0004 (7) | 0.0042 (7) | -0.0006 (7) |
| C16 | 0.0151 (8) | 0.0152 (8) | 0.0171 (9) | 0.0007 (6) | 0.0011 (7) | -0.0007 (6) |
| C17 | 0.0174 (9) | 0.0153 (8) | 0.0149 (8) | 0.0005 (6) | 0.0029 (7) | 0.0006 (6) |
| C18 | 0.0199 (9) | 0.0216 (9) | 0.0169 (9) | -0.0035 (7) | 0.0025 (7) | -0.0006 (7) |
| C19 | 0.0188 (9) | 0.0226 (9) | 0.0176 (9) | -0.0004 (7) | 0.0008 (7) | -0.0006 (7) |
| C20 | 0.0179 (9) | 0.0147 (8) | 0.0149 (9) | 0.0015 (7) | 0.0004 (7) | -0.0002 (6) |
| C20A | 0.0284 (10) | 0.0172 (8) | 0.0210 (9) | 0.0051 (7) | -0.0033 (7) | -0.0003 (7) |
| C21 | 0.0181 (9) | 0.0171 (8) | 0.0149 (8) | 0.0028 (7) | 0.0031 (7) | 0.0004 (6) |
| C22 | 0.0212 (9) | 0.0288 (10) | 0.0183 (9) | -0.0005 (7) | 0.0046 (7) | -0.0043 (7) |
| C23 | 0.0228 (9) | 0.0242 (9) | 0.0172 (9) | 0.0019 (7) | 0.0040 (7) | -0.0014 (7) |
| C24 | 0.0184 (9) | 0.0174 (9) | 0.0159 (9) | 0.0041 (7) | 0.0009 (7) | 0.0001 (7) |
| C25 | 0.0217 (9) | 0.0216 (9) | 0.0184 (9) | 0.0015 (7) | 0.0015 (7) | -0.0034 (7) |
| C25A | 0.0263 (11) | 0.0478 (13) | 0.0241 (11) | -0.0071 (9) | 0.0037 (8) | -0.0120 (9) |
| C26 | 0.0203 (9) | 0.0213 (9) | 0.0193 (9) | 0.0044 (7) | -0.0033 (7) | -0.0012 (7) |
| C27 | 0.0191 (9) | 0.0178 (9) | 0.0178 (9) | 0.0015 (7) | -0.0011 (7) | 0.0007 (7) |
| C28 | 0.0227 (9) | 0.0193 (8) | 0.0178 (9) | 0.0055 (7) | 0.0019 (7) | -0.0007 (7) |
| C29 | 0.0235 (9) | 0.0222 (9) | 0.0178 (9) | 0.0015 (7) | 0.0021 (7) | 0.0002 (7) |
| C30 | 0.0290 (11) | 0.0367 (11) | 0.0269 (11) | -0.0078 (9) | 0.0038 (8) | -0.0029 (9) |
| C31 | 0.0363 (12) | 0.0353 (11) | 0.0225 (10) | -0.0018 (9) | 0.0065 (9) | 0.0018 (8) |
| C32 | 0.0286 (11) | 0.0327 (11) | 0.0194 (10) | 0.0001 (8) | 0.0027 (8) | -0.0040 (8) |
| C33 | 0.0288 (12) | 0.0666 (17) | 0.0329 (13) | -0.0074 (11) | 0.0047 (10) | -0.0145 (11) |
| S2 | 0.0179 (2) | 0.0153 (2) | 0.0151 (2) | 0.00321 (16) | 0.00275 (16) | -0.00007 (16) |
| O4 | 0.0229 (7) | 0.0170 (6) | 0.0137 (6) | 0.0044 (5) | 0.0004 (5) | -0.0008 (5) |
| O5 | 0.0204 (7) | 0.0202 (7) | 0.0262 (8) | 0.0048 (5) | 0.0056 (6) | 0.0024 (6) |
| O6 | 0.0286 (7) | 0.0208 (7) | 0.0151 (7) | 0.0042 (6) | 0.0031 (5) | -0.0006 (5) |
| C34 | 0.0282 (11) | 0.0248 (10) | 0.0256 (11) | 0.0006 (8) | 0.0090 (8) | 0.0007 (8) |
| C35 | 0.0239 (10) | 0.0187 (9) | 0.0175 (9) | 0.0008 (7) | 0.0046 (7) | -0.0046 (7) |
| C36 | 0.0256 (10) | 0.0189 (9) | 0.0144 (9) | 0.0045 (7) | 0.0001 (7) | 0.0000 (7) |
| C37 | 0.0185 (9) | 0.0198 (9) | 0.0160 (9) | 0.0051 (7) | -0.0003 (7) | -0.0033 (7) |
| C38 | 0.0201 (9) | 0.0137 (8) | 0.0145 (9) | 0.0010 (7) | 0.0033 (7) | -0.0018 (7) |
| C39 | 0.0206 (9) | 0.0179 (9) | 0.0198 (9) | 0.0065 (7) | 0.0008 (7) | -0.0001 (7) |
| C40 | 0.0175 (9) | 0.0216 (9) | 0.0246 (10) | 0.0038 (7) | 0.0031 (7) | -0.0032 (7) |
| C41 | 0.0195 (9) | 0.0169 (9) | 0.0135 (9) | 0.0028 (7) | -0.0018 (7) | -0.0004 (7) |
| C42 | 0.0167 (9) | 0.0183 (9) | 0.0170 (9) | -0.0006 (7) | 0.0019 (7) | -0.0013 (7) |
| C43 | 0.0145 (9) | 0.0194 (9) | 0.0185 (9) | -0.0001 (7) | 0.0019 (7) | -0.0015 (7) |
| C44 | 0.0156 (8) | 0.0173 (8) | 0.0131 (8) | 0.0031 (7) | 0.0006 (6) | -0.0005 (6) |
| C44A | 0.0183 (9) | 0.0186 (8) | 0.0178 (9) | 0.0050 (7) | -0.0016 (7) | -0.0005 (7) |
| C45 | 0.0175 (9) | 0.0144 (8) | 0.0158 (9) | 0.0012 (6) | 0.0040 (7) | -0.0003 (6) |
| C46 | 0.0191 (9) | 0.0198 (9) | 0.0185 (9) | 0.0007 (7) | 0.0051 (7) | 0.0017 (7) |
| C47 | 0.0154 (8) | 0.0242 (9) | 0.0190 (9) | 0.0022 (7) | 0.0061 (7) | -0.0020 (7) |
| C48 | 0.0131 (8) | 0.0319 (10) | 0.0207 (9) | 0.0010 (7) | 0.0023 (7) | -0.0028 (7) |
| C49 | 0.0119 (8) | 0.0222 (9) | 0.0159 (9) | 0.0024 (7) | 0.0011 (6) | -0.0007 (7) |
| C50 | 0.0134 (8) | 0.0159 (8) | 0.0154 (9) | 0.0015 (6) | -0.0001 (6) | 0.0000 (6) |
| C51 | 0.0136 (9) | 0.0252 (9) | 0.0175 (9) | -0.0019 (7) | 0.0019 (7) | -0.0019 (7) |

| | | | | | | |
|------|-------------|-------------|-------------|-------------|-------------|--------------|
| C52 | 0.0200 (9) | 0.0199 (9) | 0.0158 (9) | -0.0012 (7) | 0.0032 (7) | -0.0015 (7) |
| C53 | 0.0139 (8) | 0.0166 (8) | 0.0143 (8) | 0.0010 (6) | -0.0007 (6) | 0.0010 (6) |
| C53A | 0.0207 (9) | 0.0207 (9) | 0.0187 (9) | 0.0071 (7) | 0.0006 (7) | 0.0013 (7) |
| C54 | 0.0133 (8) | 0.0206 (9) | 0.0178 (9) | 0.0034 (7) | -0.0012 (7) | -0.0009 (7) |
| C55 | 0.0152 (9) | 0.0408 (11) | 0.0201 (9) | -0.0015 (8) | -0.0006 (7) | -0.0052 (8) |
| C56 | 0.0163 (9) | 0.0367 (11) | 0.0209 (10) | 0.0025 (8) | -0.0019 (7) | -0.0050 (8) |
| C57 | 0.0181 (9) | 0.0197 (9) | 0.0154 (9) | 0.0054 (7) | -0.0024 (7) | -0.0019 (7) |
| C58 | 0.0214 (9) | 0.0181 (8) | 0.0167 (9) | 0.0067 (7) | -0.0005 (7) | -0.0026 (6) |
| C58A | 0.0224 (10) | 0.0318 (10) | 0.0189 (9) | 0.0017 (8) | 0.0033 (7) | -0.0055 (7) |
| C59 | 0.0237 (9) | 0.0181 (8) | 0.0173 (9) | 0.0072 (7) | -0.0001 (7) | -0.0002 (7) |
| C60 | 0.0185 (9) | 0.0175 (8) | 0.0204 (9) | 0.0045 (7) | -0.0002 (7) | 0.0000 (7) |
| C61 | 0.0218 (9) | 0.0193 (8) | 0.0193 (9) | 0.0029 (7) | -0.0030 (7) | -0.0028 (7) |
| C62 | 0.0197 (9) | 0.0284 (10) | 0.0262 (10) | 0.0036 (8) | -0.0006 (7) | -0.0034 (8) |
| C63 | 0.0240 (11) | 0.0482 (13) | 0.0390 (13) | 0.0088 (9) | 0.0057 (9) | -0.0063 (10) |
| C64 | 0.0259 (11) | 0.0259 (10) | 0.0538 (14) | -0.0008 (8) | 0.0049 (9) | 0.0004 (9) |
| C65 | 0.0276 (11) | 0.0337 (11) | 0.0198 (10) | -0.0015 (8) | -0.0007 (8) | -0.0072 (8) |
| C66 | 0.0340 (12) | 0.0375 (12) | 0.0263 (11) | -0.0018 (9) | 0.0048 (9) | 0.0044 (9) |

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

| | | | |
|----------|-------------|----------|-------------|
| S1—O2 | 1.4257 (13) | S2—O6 | 1.4292 (14) |
| S1—O3 | 1.4315 (13) | S2—O5 | 1.4304 (14) |
| S1—O1 | 1.5674 (13) | S2—O4 | 1.5671 (13) |
| S1—C5 | 1.7593 (18) | S2—C38 | 1.7617 (19) |
| O1—C8 | 1.479 (2) | O4—C41 | 1.483 (2) |
| C1—C2 | 1.504 (3) | C34—C35 | 1.507 (3) |
| C1—H1A | 0.9800 | C34—H34A | 0.9800 |
| C1—H1B | 0.9800 | C34—H34B | 0.9800 |
| C1—H1C | 0.9800 | C34—H34C | 0.9800 |
| C2—C3 | 1.392 (3) | C35—C40 | 1.393 (3) |
| C2—C7 | 1.398 (3) | C35—C36 | 1.394 (3) |
| C3—C4 | 1.389 (3) | C36—C37 | 1.390 (3) |
| C3—H3 | 0.9500 | C36—H36 | 0.9500 |
| C4—C5 | 1.386 (2) | C37—C38 | 1.386 (3) |
| C4—H4 | 0.9500 | C37—H37 | 0.9500 |
| C5—C6 | 1.390 (3) | C38—C39 | 1.389 (3) |
| C6—C7 | 1.382 (3) | C39—C40 | 1.394 (3) |
| C6—H6 | 0.9500 | C39—H39 | 0.9500 |
| C7—H7 | 0.9500 | C40—H40 | 0.9500 |
| C8—C9 | 1.510 (3) | C41—C42 | 1.505 (3) |
| C8—C13 | 1.519 (3) | C41—C46 | 1.518 (3) |
| C8—H8 | 1.0000 | C41—H41 | 1.0000 |
| C9—C10 | 1.530 (3) | C42—C43 | 1.530 (2) |
| C9—H9A | 0.9900 | C42—H42A | 0.9900 |
| C9—H9B | 0.9900 | C42—H42B | 0.9900 |
| C10—C11 | 1.549 (2) | C43—C44 | 1.541 (2) |
| C10—H10A | 0.9900 | C43—H43A | 0.9900 |
| C10—H10B | 0.9900 | C43—H43B | 0.9900 |

| | | | |
|-----------|-----------|-----------|-----------|
| C11—C12 | 1.526 (2) | C44—C45 | 1.528 (2) |
| C11—C11A | 1.544 (2) | C44—C44A | 1.547 (2) |
| C11—C17 | 1.556 (2) | C44—C50 | 1.555 (2) |
| C11A—H11A | 0.9800 | C44A—H44A | 0.9800 |
| C11A—H11B | 0.9800 | C44A—H44B | 0.9800 |
| C11A—H11C | 0.9800 | C44A—H44C | 0.9800 |
| C12—C14 | 1.335 (2) | C45—C47 | 1.331 (2) |
| C12—C13 | 1.519 (3) | C45—C46 | 1.517 (3) |
| C13—H13A | 0.9900 | C46—H46A | 0.9900 |
| C13—H13B | 0.9900 | C46—H46B | 0.9900 |
| C14—C15 | 1.502 (2) | C47—C48 | 1.503 (2) |
| C14—H14 | 0.9500 | C47—H47 | 0.9500 |
| C15—C16 | 1.536 (2) | C48—C49 | 1.534 (2) |
| C15—H15A | 0.9900 | C48—H48A | 0.9900 |
| C15—H15B | 0.9900 | C48—H48B | 0.9900 |
| C16—C21 | 1.523 (2) | C49—C54 | 1.528 (2) |
| C16—C17 | 1.537 (2) | C49—C50 | 1.539 (2) |
| C16—H16 | 1.0000 | C49—H49 | 1.0000 |
| C17—C18 | 1.538 (2) | C50—C51 | 1.536 (2) |
| C17—H17 | 1.0000 | C50—H50 | 1.0000 |
| C18—C19 | 1.538 (2) | C51—C52 | 1.542 (2) |
| C18—H18A | 0.9900 | C51—H51A | 0.9900 |
| C18—H18B | 0.9900 | C51—H51B | 0.9900 |
| C19—C20 | 1.532 (2) | C52—C53 | 1.535 (2) |
| C19—H19A | 0.9900 | C52—H52A | 0.9900 |
| C19—H19B | 0.9900 | C52—H52B | 0.9900 |
| C20—C20A | 1.536 (2) | C53—C53A | 1.542 (2) |
| C20—C21 | 1.547 (2) | C53—C54 | 1.545 (2) |
| C20—C24 | 1.557 (2) | C53—C57 | 1.552 (2) |
| C20A—H20A | 0.9800 | C53A—H53A | 0.9800 |
| C20A—H20B | 0.9800 | C53A—H53B | 0.9800 |
| C20A—H20C | 0.9800 | C53A—H53C | 0.9800 |
| C21—C22 | 1.527 (2) | C54—C55 | 1.533 (3) |
| C21—H21 | 1.0000 | C54—H54 | 1.0000 |
| C22—C23 | 1.546 (3) | C55—C56 | 1.550 (3) |
| C22—H22A | 0.9900 | C55—H55A | 0.9900 |
| C22—H22B | 0.9900 | C55—H55B | 0.9900 |
| C23—C24 | 1.556 (3) | C56—C57 | 1.552 (3) |
| C23—H23A | 0.9900 | C56—H56A | 0.9900 |
| C23—H23B | 0.9900 | C56—H56B | 0.9900 |
| C24—C25 | 1.543 (3) | C57—C58 | 1.542 (2) |
| C24—H24 | 1.0000 | C57—H57 | 1.0000 |
| C25—C26 | 1.508 (3) | C58—C59 | 1.508 (2) |
| C25—C25A | 1.539 (3) | C58—C58A | 1.533 (3) |
| C25—H25 | 1.0000 | C58—H58 | 1.0000 |
| C25A—H25A | 0.9800 | C58A—H58A | 0.9800 |
| C25A—H25B | 0.9800 | C58A—H58B | 0.9800 |
| C25A—H25C | 0.9800 | C58A—H58C | 0.9800 |

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|------------|-------------|---------------|-------------|
| C26—C27 | 1.322 (3) | C59—C60 | 1.324 (2) |
| C26—H26 | 0.9500 | C59—H59 | 0.9500 |
| C27—C28 | 1.503 (2) | C60—C61 | 1.506 (2) |
| C27—H27 | 0.9500 | C60—H60 | 0.9500 |
| C28—C29 | 1.542 (3) | C61—C65 | 1.536 (3) |
| C28—C32 | 1.544 (3) | C61—C62 | 1.541 (3) |
| C28—H28 | 1.0000 | C61—H61 | 1.0000 |
| C29—C30 | 1.531 (3) | C62—C64 | 1.522 (3) |
| C29—C31 | 1.534 (3) | C62—C63 | 1.523 (3) |
| C29—H29 | 1.0000 | C62—H62 | 1.0000 |
| C30—H30A | 0.9800 | C63—H63A | 0.9800 |
| C30—H30B | 0.9800 | C63—H63B | 0.9800 |
| C30—H30C | 0.9800 | C63—H63C | 0.9800 |
| C31—H31A | 0.9800 | C64—H64A | 0.9800 |
| C31—H31B | 0.9800 | C64—H64B | 0.9800 |
| C31—H31C | 0.9800 | C64—H64C | 0.9800 |
| C32—C33 | 1.518 (3) | C65—C66 | 1.518 (3) |
| C32—H32A | 0.9900 | C65—H65A | 0.9900 |
| C32—H32B | 0.9900 | C65—H65B | 0.9900 |
| C33—H33A | 0.9800 | C66—H66A | 0.9800 |
| C33—H33B | 0.9800 | C66—H66B | 0.9800 |
| C33—H33C | 0.9800 | C66—H66C | 0.9800 |
| | | | |
| O2—S1—O3 | 119.33 (8) | O6—S2—O5 | 119.26 (8) |
| O2—S1—O1 | 104.43 (7) | O6—S2—O4 | 104.35 (7) |
| O3—S1—O1 | 110.39 (8) | O5—S2—O4 | 109.84 (8) |
| O2—S1—C5 | 109.95 (8) | O6—S2—C38 | 109.82 (9) |
| O3—S1—C5 | 108.60 (8) | O5—S2—C38 | 108.38 (8) |
| O1—S1—C5 | 102.86 (8) | O4—S2—C38 | 104.11 (8) |
| C8—O1—S1 | 117.46 (11) | C41—O4—S2 | 118.40 (11) |
| C2—C1—H1A | 109.5 | C35—C34—H34A | 109.5 |
| C2—C1—H1B | 109.5 | C35—C34—H34B | 109.5 |
| H1A—C1—H1B | 109.5 | H34A—C34—H34B | 109.5 |
| C2—C1—H1C | 109.5 | C35—C34—H34C | 109.5 |
| H1A—C1—H1C | 109.5 | H34A—C34—H34C | 109.5 |
| H1B—C1—H1C | 109.5 | H34B—C34—H34C | 109.5 |
| C3—C2—C7 | 118.71 (17) | C40—C35—C36 | 118.45 (18) |
| C3—C2—C1 | 120.74 (18) | C40—C35—C34 | 120.45 (18) |
| C7—C2—C1 | 120.55 (18) | C36—C35—C34 | 121.08 (18) |
| C4—C3—C2 | 120.85 (17) | C37—C36—C35 | 121.01 (17) |
| C4—C3—H3 | 119.6 | C37—C36—H36 | 119.5 |
| C2—C3—H3 | 119.6 | C35—C36—H36 | 119.5 |
| C5—C4—C3 | 119.17 (17) | C38—C37—C36 | 119.37 (17) |
| C5—C4—H4 | 120.4 | C38—C37—H37 | 120.3 |
| C3—C4—H4 | 120.4 | C36—C37—H37 | 120.3 |
| C4—C5—C6 | 121.14 (17) | C37—C38—C39 | 121.01 (17) |
| C4—C5—S1 | 120.04 (14) | C37—C38—S2 | 119.81 (14) |
| C6—C5—S1 | 118.82 (14) | C39—C38—S2 | 119.16 (14) |

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| C7—C6—C5 | 118.96 (18) | C38—C39—C40 | 118.78 (17) |
| C7—C6—H6 | 120.5 | C38—C39—H39 | 120.6 |
| C5—C6—H6 | 120.5 | C40—C39—H39 | 120.6 |
| C6—C7—C2 | 121.17 (18) | C35—C40—C39 | 121.38 (18) |
| C6—C7—H7 | 119.4 | C35—C40—H40 | 119.3 |
| C2—C7—H7 | 119.4 | C39—C40—H40 | 119.3 |
| O1—C8—C9 | 107.66 (14) | O4—C41—C42 | 108.23 (14) |
| O1—C8—C13 | 107.31 (15) | O4—C41—C46 | 107.85 (14) |
| C9—C8—C13 | 112.22 (15) | C42—C41—C46 | 111.52 (15) |
| O1—C8—H8 | 109.9 | O4—C41—H41 | 109.7 |
| C9—C8—H8 | 109.9 | C42—C41—H41 | 109.7 |
| C13—C8—H8 | 109.9 | C46—C41—H41 | 109.7 |
| C8—C9—C10 | 110.04 (15) | C41—C42—C43 | 109.48 (15) |
| C8—C9—H9A | 109.7 | C41—C42—H42A | 109.8 |
| C10—C9—H9A | 109.7 | C43—C42—H42A | 109.8 |
| C8—C9—H9B | 109.7 | C41—C42—H42B | 109.8 |
| C10—C9—H9B | 109.7 | C43—C42—H42B | 109.8 |
| H9A—C9—H9B | 108.2 | H42A—C42—H42B | 108.2 |
| C9—C10—C11 | 114.88 (16) | C42—C43—C44 | 114.46 (15) |
| C9—C10—H10A | 108.5 | C42—C43—H43A | 108.6 |
| C11—C10—H10A | 108.5 | C44—C43—H43A | 108.6 |
| C9—C10—H10B | 108.5 | C42—C43—H43B | 108.6 |
| C11—C10—H10B | 108.5 | C44—C43—H43B | 108.6 |
| H10A—C10—H10B | 107.5 | H43A—C43—H43B | 107.6 |
| C12—C11—C11A | 107.17 (15) | C45—C44—C43 | 109.69 (14) |
| C12—C11—C10 | 110.18 (14) | C45—C44—C44A | 107.78 (14) |
| C11A—C11—C10 | 109.72 (15) | C43—C44—C44A | 109.68 (15) |
| C12—C11—C17 | 108.98 (14) | C45—C44—C50 | 108.97 (14) |
| C11A—C11—C17 | 112.24 (14) | C43—C44—C50 | 108.65 (14) |
| C10—C11—C17 | 108.54 (14) | C44A—C44—C50 | 112.04 (14) |
| C11—C11A—H11A | 109.5 | C44—C44A—H44A | 109.5 |
| C11—C11A—H11B | 109.5 | C44—C44A—H44B | 109.5 |
| H11A—C11A—H11B | 109.5 | H44A—C44A—H44B | 109.5 |
| C11—C11A—H11C | 109.5 | C44—C44A—H44C | 109.5 |
| H11A—C11A—H11C | 109.5 | H44A—C44A—H44C | 109.5 |
| H11B—C11A—H11C | 109.5 | H44B—C44A—H44C | 109.5 |
| C14—C12—C13 | 120.64 (16) | C47—C45—C46 | 120.50 (16) |
| C14—C12—C11 | 122.48 (16) | C47—C45—C44 | 122.27 (16) |
| C13—C12—C11 | 116.71 (15) | C46—C45—C44 | 117.15 (15) |
| C8—C13—C12 | 110.98 (15) | C45—C46—C41 | 111.39 (15) |
| C8—C13—H13A | 109.4 | C45—C46—H46A | 109.3 |
| C12—C13—H13A | 109.4 | C41—C46—H46A | 109.3 |
| C8—C13—H13B | 109.4 | C45—C46—H46B | 109.3 |
| C12—C13—H13B | 109.4 | C41—C46—H46B | 109.3 |
| H13A—C13—H13B | 108.0 | H46A—C46—H46B | 108.0 |
| C12—C14—C15 | 124.96 (16) | C45—C47—C48 | 125.15 (16) |
| C12—C14—H14 | 117.5 | C45—C47—H47 | 117.4 |
| C15—C14—H14 | 117.5 | C48—C47—H47 | 117.4 |

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| C14—C15—C16 | 113.52 (14) | C47—C48—C49 | 113.49 (14) |
| C14—C15—H15A | 108.9 | C47—C48—H48A | 108.9 |
| C16—C15—H15A | 108.9 | C49—C48—H48A | 108.9 |
| C14—C15—H15B | 108.9 | C47—C48—H48B | 108.9 |
| C16—C15—H15B | 108.9 | C49—C48—H48B | 108.9 |
| H15A—C15—H15B | 107.7 | H48A—C48—H48B | 107.7 |
| C21—C16—C15 | 111.21 (14) | C54—C49—C48 | 111.06 (14) |
| C21—C16—C17 | 110.09 (14) | C54—C49—C50 | 110.31 (14) |
| C15—C16—C17 | 109.65 (14) | C48—C49—C50 | 109.12 (14) |
| C21—C16—H16 | 108.6 | C54—C49—H49 | 108.8 |
| C15—C16—H16 | 108.6 | C48—C49—H49 | 108.8 |
| C17—C16—H16 | 108.6 | C50—C49—H49 | 108.8 |
| C16—C17—C18 | 113.36 (14) | C51—C50—C49 | 112.82 (14) |
| C16—C17—C11 | 111.53 (14) | C51—C50—C44 | 112.92 (14) |
| C18—C17—C11 | 113.09 (14) | C49—C50—C44 | 111.62 (14) |
| C16—C17—H17 | 106.1 | C51—C50—H50 | 106.3 |
| C18—C17—H17 | 106.1 | C49—C50—H50 | 106.3 |
| C11—C17—H17 | 106.1 | C44—C50—H50 | 106.3 |
| C17—C18—C19 | 114.33 (15) | C50—C51—C52 | 114.02 (14) |
| C17—C18—H18A | 108.7 | C50—C51—H51A | 108.7 |
| C19—C18—H18A | 108.7 | C52—C51—H51A | 108.7 |
| C17—C18—H18B | 108.7 | C50—C51—H51B | 108.7 |
| C19—C18—H18B | 108.7 | C52—C51—H51B | 108.7 |
| H18A—C18—H18B | 107.6 | H51A—C51—H51B | 107.6 |
| C20—C19—C18 | 111.48 (15) | C53—C52—C51 | 110.86 (14) |
| C20—C19—H19A | 109.3 | C53—C52—H52A | 109.5 |
| C18—C19—H19A | 109.3 | C51—C52—H52A | 109.5 |
| C20—C19—H19B | 109.3 | C53—C52—H52B | 109.5 |
| C18—C19—H19B | 109.3 | C51—C52—H52B | 109.5 |
| H19A—C19—H19B | 108.0 | H52A—C52—H52B | 108.1 |
| C19—C20—C20A | 110.43 (15) | C52—C53—C53A | 110.86 (14) |
| C19—C20—C21 | 106.59 (14) | C52—C53—C54 | 106.80 (14) |
| C20A—C20—C21 | 112.48 (15) | C53A—C53—C54 | 112.19 (14) |
| C19—C20—C24 | 116.53 (14) | C52—C53—C57 | 116.91 (14) |
| C20A—C20—C24 | 110.01 (14) | C53A—C53—C57 | 109.47 (14) |
| C21—C20—C24 | 100.44 (13) | C54—C53—C57 | 100.16 (13) |
| C20—C20A—H20A | 109.5 | C53—C53A—H53A | 109.5 |
| C20—C20A—H20B | 109.5 | C53—C53A—H53B | 109.5 |
| H20A—C20A—H20B | 109.5 | H53A—C53A—H53B | 109.5 |
| C20—C20A—H20C | 109.5 | C53—C53A—H53C | 109.5 |
| H20A—C20A—H20C | 109.5 | H53A—C53A—H53C | 109.5 |
| H20B—C20A—H20C | 109.5 | H53B—C53A—H53C | 109.5 |
| C16—C21—C22 | 118.58 (15) | C49—C54—C55 | 118.65 (15) |
| C16—C21—C20 | 114.46 (14) | C49—C54—C53 | 114.87 (14) |
| C22—C21—C20 | 104.56 (14) | C55—C54—C53 | 104.15 (14) |
| C16—C21—H21 | 106.1 | C49—C54—H54 | 106.1 |
| C22—C21—H21 | 106.1 | C55—C54—H54 | 106.1 |
| C20—C21—H21 | 106.1 | C53—C54—H54 | 106.1 |

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| C21—C22—C23 | 104.07 (14) | C54—C55—C56 | 104.04 (15) |
| C21—C22—H22A | 110.9 | C54—C55—H55A | 110.9 |
| C23—C22—H22A | 110.9 | C56—C55—H55A | 110.9 |
| C21—C22—H22B | 110.9 | C54—C55—H55B | 110.9 |
| C23—C22—H22B | 110.9 | C56—C55—H55B | 110.9 |
| H22A—C22—H22B | 109.0 | H55A—C55—H55B | 109.0 |
| C22—C23—C24 | 107.21 (14) | C55—C56—C57 | 106.77 (14) |
| C22—C23—H23A | 110.3 | C55—C56—H56A | 110.4 |
| C24—C23—H23A | 110.3 | C57—C56—H56A | 110.4 |
| C22—C23—H23B | 110.3 | C55—C56—H56B | 110.4 |
| C24—C23—H23B | 110.3 | C57—C56—H56B | 110.4 |
| H23A—C23—H23B | 108.5 | H56A—C56—H56B | 108.6 |
| C25—C24—C23 | 111.24 (14) | C58—C57—C56 | 112.65 (15) |
| C25—C24—C20 | 118.58 (15) | C58—C57—C53 | 118.51 (14) |
| C23—C24—C20 | 103.41 (14) | C56—C57—C53 | 103.44 (14) |
| C25—C24—H24 | 107.7 | C58—C57—H57 | 107.2 |
| C23—C24—H24 | 107.7 | C56—C57—H57 | 107.2 |
| C20—C24—H24 | 107.7 | C53—C57—H57 | 107.2 |
| C26—C25—C25A | 109.16 (16) | C59—C58—C58A | 108.23 (15) |
| C26—C25—C24 | 110.43 (15) | C59—C58—C57 | 111.30 (14) |
| C25A—C25—C24 | 113.53 (15) | C58A—C58—C57 | 112.29 (14) |
| C26—C25—H25 | 107.8 | C59—C58—H58 | 108.3 |
| C25A—C25—H25 | 107.8 | C58A—C58—H58 | 108.3 |
| C24—C25—H25 | 107.8 | C57—C58—H58 | 108.3 |
| C25—C25A—H25A | 109.5 | C58—C58A—H58A | 109.5 |
| C25—C25A—H25B | 109.5 | C58—C58A—H58B | 109.5 |
| H25A—C25A—H25B | 109.5 | H58A—C58A—H58B | 109.5 |
| C25—C25A—H25C | 109.5 | C58—C58A—H58C | 109.5 |
| H25A—C25A—H25C | 109.5 | H58A—C58A—H58C | 109.5 |
| H25B—C25A—H25C | 109.5 | H58B—C58A—H58C | 109.5 |
| C27—C26—C25 | 124.38 (17) | C60—C59—C58 | 125.96 (16) |
| C27—C26—H26 | 117.8 | C60—C59—H59 | 117.0 |
| C25—C26—H26 | 117.8 | C58—C59—H59 | 117.0 |
| C26—C27—C28 | 126.95 (17) | C59—C60—C61 | 125.34 (16) |
| C26—C27—H27 | 116.5 | C59—C60—H60 | 117.3 |
| C28—C27—H27 | 116.5 | C61—C60—H60 | 117.3 |
| C27—C28—C29 | 111.71 (15) | C60—C61—C65 | 109.78 (15) |
| C27—C28—C32 | 108.93 (15) | C60—C61—C62 | 113.62 (16) |
| C29—C28—C32 | 111.10 (15) | C65—C61—C62 | 111.72 (15) |
| C27—C28—H28 | 108.3 | C60—C61—H61 | 107.1 |
| C29—C28—H28 | 108.3 | C65—C61—H61 | 107.1 |
| C32—C28—H28 | 108.3 | C62—C61—H61 | 107.1 |
| C30—C29—C31 | 109.07 (16) | C64—C62—C63 | 110.29 (18) |
| C30—C29—C28 | 112.32 (15) | C64—C62—C61 | 113.65 (16) |
| C31—C29—C28 | 112.07 (16) | C63—C62—C61 | 111.00 (16) |
| C30—C29—H29 | 107.7 | C64—C62—H62 | 107.2 |
| C31—C29—H29 | 107.7 | C63—C62—H62 | 107.2 |
| C28—C29—H29 | 107.7 | C61—C62—H62 | 107.2 |

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| C29—C30—H30A | 109.5 | C62—C63—H63A | 109.5 |
| C29—C30—H30B | 109.5 | C62—C63—H63B | 109.5 |
| H30A—C30—H30B | 109.5 | H63A—C63—H63B | 109.5 |
| C29—C30—H30C | 109.5 | C62—C63—H63C | 109.5 |
| H30A—C30—H30C | 109.5 | H63A—C63—H63C | 109.5 |
| H30B—C30—H30C | 109.5 | H63B—C63—H63C | 109.5 |
| C29—C31—H31A | 109.5 | C62—C64—H64A | 109.5 |
| C29—C31—H31B | 109.5 | C62—C64—H64B | 109.5 |
| H31A—C31—H31B | 109.5 | H64A—C64—H64B | 109.5 |
| C29—C31—H31C | 109.5 | C62—C64—H64C | 109.5 |
| H31A—C31—H31C | 109.5 | H64A—C64—H64C | 109.5 |
| H31B—C31—H31C | 109.5 | H64B—C64—H64C | 109.5 |
| C33—C32—C28 | 114.20 (17) | C66—C65—C61 | 113.44 (16) |
| C33—C32—H32A | 108.7 | C66—C65—H65A | 108.9 |
| C28—C32—H32A | 108.7 | C61—C65—H65A | 108.9 |
| C33—C32—H32B | 108.7 | C66—C65—H65B | 108.9 |
| C28—C32—H32B | 108.7 | C61—C65—H65B | 108.9 |
| H32A—C32—H32B | 107.6 | H65A—C65—H65B | 107.7 |
| C32—C33—H33A | 109.5 | C65—C66—H66A | 109.5 |
| C32—C33—H33B | 109.5 | C65—C66—H66B | 109.5 |
| H33A—C33—H33B | 109.5 | H66A—C66—H66B | 109.5 |
| C32—C33—H33C | 109.5 | C65—C66—H66C | 109.5 |
| H33A—C33—H33C | 109.5 | H66A—C66—H66C | 109.5 |
| H33B—C33—H33C | 109.5 | H66B—C66—H66C | 109.5 |
| | | | |
| O2—S1—O1—C8 | -177.24 (12) | O6—S2—O4—C41 | -178.70 (12) |
| O3—S1—O1—C8 | -47.79 (14) | O5—S2—O4—C41 | 52.34 (14) |
| C5—S1—O1—C8 | 67.92 (13) | C38—S2—O4—C41 | -63.53 (14) |
| C7—C2—C3—C4 | 0.4 (3) | C40—C35—C36—C37 | -0.8 (3) |
| C1—C2—C3—C4 | -178.74 (18) | C34—C35—C36—C37 | 177.58 (17) |
| C2—C3—C4—C5 | -0.1 (3) | C35—C36—C37—C38 | 0.2 (3) |
| C3—C4—C5—C6 | -0.4 (3) | C36—C37—C38—C39 | 0.6 (3) |
| C3—C4—C5—S1 | 179.51 (14) | C36—C37—C38—S2 | 179.16 (14) |
| O2—S1—C5—C4 | 129.06 (15) | O6—S2—C38—C37 | -132.77 (15) |
| O3—S1—C5—C4 | -3.17 (17) | O5—S2—C38—C37 | -0.91 (17) |
| O1—S1—C5—C4 | -120.16 (15) | O4—S2—C38—C37 | 115.98 (15) |
| O2—S1—C5—C6 | -51.05 (17) | O6—S2—C38—C39 | 45.84 (16) |
| O3—S1—C5—C6 | 176.72 (14) | O5—S2—C38—C39 | 177.70 (14) |
| O1—S1—C5—C6 | 59.72 (16) | O4—S2—C38—C39 | -65.41 (16) |
| C4—C5—C6—C7 | 0.4 (3) | C37—C38—C39—C40 | -0.7 (3) |
| S1—C5—C6—C7 | -179.44 (15) | S2—C38—C39—C40 | -179.27 (14) |
| C5—C6—C7—C2 | -0.1 (3) | C36—C35—C40—C39 | 0.7 (3) |
| C3—C2—C7—C6 | -0.4 (3) | C34—C35—C40—C39 | -177.70 (17) |
| C1—C2—C7—C6 | 178.82 (19) | C38—C39—C40—C35 | 0.1 (3) |
| S1—O1—C8—C9 | 113.97 (14) | S2—O4—C41—C42 | 116.19 (14) |
| S1—O1—C8—C13 | -125.04 (13) | S2—O4—C41—C46 | -123.02 (13) |
| O1—C8—C9—C10 | 175.44 (15) | O4—C41—C42—C43 | 177.69 (14) |
| C13—C8—C9—C10 | 57.6 (2) | C46—C41—C42—C43 | 59.2 (2) |

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| C8—C9—C10—C11 | −54.0 (2) | C41—C42—C43—C44 | −57.0 (2) |
| C9—C10—C11—C12 | 46.7 (2) | C42—C43—C44—C45 | 48.1 (2) |
| C9—C10—C11—C11A | −71.1 (2) | C42—C43—C44—C44A | −70.12 (19) |
| C9—C10—C11—C17 | 165.97 (15) | C42—C43—C44—C50 | 167.10 (14) |
| C11A—C11—C12—C14 | −100.7 (2) | C43—C44—C45—C47 | 139.53 (17) |
| C10—C11—C12—C14 | 140.00 (18) | C44A—C44—C45—C47 | −101.10 (19) |
| C17—C11—C12—C14 | 21.0 (2) | C50—C44—C45—C47 | 20.7 (2) |
| C11A—C11—C12—C13 | 74.55 (19) | C43—C44—C45—C46 | −43.9 (2) |
| C10—C11—C12—C13 | −44.8 (2) | C44A—C44—C45—C46 | 75.46 (19) |
| C17—C11—C12—C13 | −163.77 (15) | C50—C44—C45—C46 | −162.74 (14) |
| O1—C8—C13—C12 | −173.35 (14) | C47—C45—C46—C41 | −135.25 (18) |
| C9—C8—C13—C12 | −55.3 (2) | C44—C45—C46—C41 | 48.1 (2) |
| C14—C12—C13—C8 | −134.95 (18) | O4—C41—C46—C45 | −173.52 (14) |
| C11—C12—C13—C8 | 49.7 (2) | C42—C41—C46—C45 | −54.8 (2) |
| C13—C12—C14—C15 | −174.45 (16) | C46—C45—C47—C48 | −176.09 (17) |
| C11—C12—C14—C15 | 0.6 (3) | C44—C45—C47—C48 | 0.4 (3) |
| C12—C14—C15—C16 | 8.2 (3) | C45—C47—C48—C49 | 9.2 (3) |
| C14—C15—C16—C21 | −160.27 (14) | C47—C48—C49—C54 | −161.03 (15) |
| C14—C15—C16—C17 | −38.3 (2) | C47—C48—C49—C50 | −39.2 (2) |
| C21—C16—C17—C18 | −46.6 (2) | C54—C49—C50—C51 | −47.0 (2) |
| C15—C16—C17—C18 | −169.27 (14) | C48—C49—C50—C51 | −169.30 (15) |
| C21—C16—C17—C11 | −175.61 (13) | C54—C49—C50—C44 | −175.44 (14) |
| C15—C16—C17—C11 | 61.72 (18) | C48—C49—C50—C44 | 62.29 (18) |
| C12—C11—C17—C16 | −51.84 (18) | C45—C44—C50—C51 | 179.78 (14) |
| C11A—C11—C17—C16 | 66.72 (18) | C43—C44—C50—C51 | 60.30 (18) |
| C10—C11—C17—C16 | −171.84 (14) | C44A—C44—C50—C51 | −61.04 (19) |
| C12—C11—C17—C18 | 179.01 (15) | C45—C44—C50—C49 | −51.86 (18) |
| C11A—C11—C17—C18 | −62.44 (19) | C43—C44—C50—C49 | −171.34 (14) |
| C10—C11—C17—C18 | 59.00 (18) | C44A—C44—C50—C49 | 67.32 (18) |
| C16—C17—C18—C19 | 46.6 (2) | C49—C50—C51—C52 | 48.9 (2) |
| C11—C17—C18—C19 | 174.85 (15) | C44—C50—C51—C52 | 176.59 (14) |
| C17—C18—C19—C20 | −52.9 (2) | C50—C51—C52—C53 | −55.0 (2) |
| C18—C19—C20—C20A | −65.06 (19) | C51—C52—C53—C53A | −64.76 (18) |
| C18—C19—C20—C21 | 57.39 (18) | C51—C52—C53—C54 | 57.74 (18) |
| C18—C19—C20—C24 | 168.49 (15) | C51—C52—C53—C57 | 168.83 (14) |
| C15—C16—C21—C22 | −57.8 (2) | C48—C49—C54—C55 | −59.8 (2) |
| C17—C16—C21—C22 | −179.55 (15) | C50—C49—C54—C55 | 179.12 (15) |
| C15—C16—C21—C20 | 177.98 (14) | C48—C49—C54—C53 | 176.18 (15) |
| C17—C16—C21—C20 | 56.24 (18) | C50—C49—C54—C53 | 55.06 (19) |
| C19—C20—C21—C16 | −61.51 (18) | C52—C53—C54—C49 | −60.29 (18) |
| C20A—C20—C21—C16 | 59.65 (19) | C53A—C53—C54—C49 | 61.37 (19) |
| C24—C20—C21—C16 | 176.56 (14) | C57—C53—C54—C49 | 177.40 (14) |
| C19—C20—C21—C22 | 167.10 (14) | C52—C53—C54—C55 | 168.28 (14) |
| C20A—C20—C21—C22 | −71.74 (18) | C53A—C53—C54—C55 | −70.06 (18) |
| C24—C20—C21—C22 | 45.18 (16) | C57—C53—C54—C55 | 45.97 (17) |
| C16—C21—C22—C23 | −161.85 (15) | C49—C54—C55—C56 | −161.69 (15) |
| C20—C21—C22—C23 | −32.90 (18) | C53—C54—C55—C56 | −32.50 (18) |
| C21—C22—C23—C24 | 7.61 (19) | C54—C55—C56—C57 | 6.2 (2) |

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| C22—C23—C24—C25 | 148.40 (15) | C55—C56—C57—C58 | 151.20 (15) |
| C22—C23—C24—C20 | 20.06 (18) | C55—C56—C57—C53 | 22.03 (18) |
| C19—C20—C24—C25 | 82.57 (19) | C52—C53—C57—C58 | 78.6 (2) |
| C20A—C20—C24—C25 | -44.1 (2) | C53A—C53—C57—C58 | -48.5 (2) |
| C21—C20—C24—C25 | -162.83 (15) | C54—C53—C57—C58 | -166.58 (15) |
| C19—C20—C24—C23 | -153.79 (15) | C52—C53—C57—C56 | -155.95 (15) |
| C20A—C20—C24—C23 | 79.55 (17) | C53A—C53—C57—C56 | 76.96 (17) |
| C21—C20—C24—C23 | -39.18 (16) | C54—C53—C57—C56 | -41.10 (16) |
| C23—C24—C25—C26 | 53.5 (2) | C56—C57—C58—C59 | 56.8 (2) |
| C20—C24—C25—C26 | 173.16 (15) | C53—C57—C58—C59 | 177.69 (15) |
| C23—C24—C25—C25A | 176.46 (17) | C56—C57—C58—C58A | 178.32 (15) |
| C20—C24—C25—C25A | -63.9 (2) | C53—C57—C58—C58A | -60.8 (2) |
| C25A—C25—C26—C27 | 117.9 (2) | C58A—C58—C59—C60 | 112.9 (2) |
| C24—C25—C26—C27 | -116.6 (2) | C57—C58—C59—C60 | -123.2 (2) |
| C25—C26—C27—C28 | -177.30 (17) | C58—C59—C60—C61 | -173.26 (17) |
| C26—C27—C28—C29 | -123.9 (2) | C59—C60—C61—C65 | 115.6 (2) |
| C26—C27—C28—C32 | 113.0 (2) | C59—C60—C61—C62 | -118.5 (2) |
| C27—C28—C29—C30 | 45.2 (2) | C60—C61—C62—C64 | -66.8 (2) |
| C32—C28—C29—C30 | 167.11 (16) | C65—C61—C62—C64 | 58.0 (2) |
| C27—C28—C29—C31 | 168.45 (16) | C60—C61—C62—C63 | 58.2 (2) |
| C32—C28—C29—C31 | -69.7 (2) | C65—C61—C62—C63 | -176.96 (17) |
| C27—C28—C32—C33 | -63.3 (2) | C60—C61—C65—C66 | -71.4 (2) |
| C29—C28—C32—C33 | 173.28 (18) | C62—C61—C65—C66 | 161.63 (17) |

Hydrogen-bond geometry (Å, °)

| D—H···A | D—H | H···A | D···A | D—H···A |
|----------------------------|------|-------|-----------|---------|
| C7—H7···O3 ⁱ | 0.95 | 2.49 | 3.217 (2) | 134 |
| C13—H13b···O6 | 0.99 | 2.54 | 3.519 (2) | 172 |
| C40—H40···O5 ⁱⁱ | 0.95 | 2.48 | 3.193 (2) | 131 |
| C42—H42a···O2 | 0.99 | 2.56 | 3.548 (2) | 175 |
| C44a—H44c···O2 | 0.98 | 2.54 | 3.390 (2) | 145 |

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