

## (*E*)-3-[4-(Dodecyloxy)phenyl]-1-(2-hydroxyphenyl)prop-2-en-1-one

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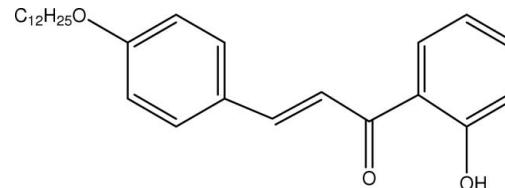
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Key indicators: single-crystal X-ray study;  $T = 100\text{ K}$ ; mean  $\sigma(\text{C}-\text{C}) = 0.005\text{ \AA}$ ;  $R$  factor = 0.069;  $wR$  factor = 0.206; data-to-parameter ratio = 15.7.

In the title compound,  $C_{27}H_{36}O_3$ , the asymmetric unit consists of two crystallographically independent molecules. The aromatic rings form dihedral angles of  $17.1(2)$  and  $17.6(2)^\circ$  in the two molecules. In both molecules, the enone groups adopt an *s-cis* conformation and the alkoxy chains are in *trans* conformations curving out of the zigzag plane. Intramolecular O—H···O hydrogen bonds involving the keto and hydroxy groups generate *S*(6) ring motifs. The molecules are stacked alternately in a head-to-tail fashion along the  $a$  axis and the crystal structure is stabilized by weak C—H···π interactions. The crystal studied was a non-merohedral twin, the ratio of components being  $0.788(2):0.212(2)$ .

### Related literature

For general background to the biological activity of chalcone derivatives, see: Bhat *et al.* (2005); Xue *et al.* (2004); Satyanarayana *et al.* (2004); Zhao *et al.* (2005); Lee *et al.* (2006). For related structures, see: Ng *et al.* (2006); Razak *et al.* (2009); Ngaini, Fadzillah *et al.* (2009); Ngaini, Rahman *et al.* (2009). For details of hydrogen-bond motifs, see: Bernstein *et al.* (1995). For stability of the temperature controller used for the data collection, see: Cosier & Glazer (1986).



### Experimental

#### Crystal data

|                              |  |
|------------------------------|--|
| $C_{27}H_{36}O_3$            | $\gamma = 84.669(5)^\circ$               |
| $M_r = 408.56$               | $V = 2302.7(3)\text{ \AA}^3$             |
| Triclinic, $P\bar{1}$        | $Z = 4$                                  |
| $a = 7.4953(6)\text{ \AA}$   | Mo $K\alpha$ radiation                   |
| $b = 13.4714(9)\text{ \AA}$  | $\mu = 0.08\text{ mm}^{-1}$              |
| $c = 23.7874(18)\text{ \AA}$ | $T = 100\text{ K}$                       |
| $\alpha = 75.116(4)^\circ$   | $0.55 \times 0.13 \times 0.06\text{ mm}$ |
| $\beta = 83.876(5)^\circ$    |  |

#### Data collection

|   |  |
|---|--|
| Bruker APEXII diffractometer                                | 8571 measured reflections              |
| Absorption correction: multi-scan<br>(SADABS; Bruker, 2005) | 8571 independent reflections           |
| $T_{\min} = 0.960$ , $T_{\max} = 0.996$                     | 4737 reflections with $I > 2\sigma(I)$ |

#### Refinement

|                                 |   |
|---------------------------------|---|
| $R[F^2 > 2\sigma(F^2)] = 0.069$ | 546 parameters                                |
| $wR(F^2) = 0.206$               | H-atom parameters constrained                 |
| $S = 1.03$                      | $\Delta\rho_{\max} = 0.29\text{ e \AA}^{-3}$  |
| 8571 reflections                | $\Delta\rho_{\min} = -0.30\text{ e \AA}^{-3}$ |

**Table 1**  
Hydrogen-bond geometry ( $\text{\AA}$ ,  $^\circ$ ).

| $D-\text{H}\cdots A$         | $D-\text{H}$ | $\text{H}\cdots A$ | $D\cdots A$ | $D-\text{H}\cdots A$ |
|------------------------------|--------------|--------------------|-------------|----------------------|
| O1A—H1A···O2A                | 0.82         | 1.79               | 2.513 (4)   | 146                  |
| O1B—H1B···O2B                | 0.82         | 1.81               | 2.530 (4)   | 146                  |
| C22B—H22C···Cg1 <sup>i</sup> | 0.97         | 2.77               | 3.654 (4)   | 151                  |
| C17B—H17D···Cg2              | 0.97         | 2.82               | 3.612 (4)   | 139                  |
| C22A—H22B···Cg3              | 0.97         | 2.93               | 3.765 (4)   | 145                  |

Symmetry code: (i)  $x - 1, y, z$ . Cg1, Cg2 and Cg3 are the centroids of the C1A–C6A, C10A–C15A and C1B–C6B rings, respectively.

Data collection: *APEX2* (Bruker, 2005); cell refinement: *SAINT* (Bruker, 2005); data reduction: *SAINT* (Bruker, 2005); program(s) used to solve structure: *SHELXTL* (Sheldrick, 2008); program(s) used to refine structure: *SHELXTL* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL* (Sheldrick, 2008).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: LH2802).

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# supporting information

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## (E)-3-[4-(Dodecyloxy)phenyl]-1-(2-hydroxyphenyl)prop-2-en-1-one

**Ibrahim Abdul Razak, Hoong-Kun Fun, Zainab Ngaini, Siti Muhaini Haris Fadzillah and Hasnain Hussain**

### S1. Comment

Chalcone derivatives are of interest because of their biological properties such as anticancer (Bhat *et al.*, 2005), antimalarial (Xue *et al.*, 2004), antiangiogenic and antitumour (Lee *et al.*, 2006), antiplatelet activity (Zhao *et al.*, 2005) and antihyperglycemic activity (Satyanarayana *et al.*, 2004). As part of our studies on their biological properties, we have synthesized the title chalcone derivative, (I). Its antibacterial activities were tested against *E. coli* ATCC 8739 and showed antimicrobial activity. The structure determination of (I) was carried out and is reported in this paper.

The asymmetric unit of (I) consists of two crystallographically independent molecules, A and B (Fig.1). The bond lengths show normal values (Allen *et al.*, 1987). The mean plane through the enone moiety (O2/C7/C8/C9) makes dihedral angles with the two benzene rings with values of 3.4 (2) $^{\circ}$  (C1—C6) and 16.0 (2) $^{\circ}$  (C10—C15) in molecule A and 7.8 (2) $^{\circ}$  (C1—C6) and 15.7 (2) $^{\circ}$  (C10—C15) in molecule B. The two benzene rings form dihedral angles with each other of 17.1 (2) $^{\circ}$  and 17.6 (2) $^{\circ}$  in molecules A and B, respectively.

The enone moieties adopt *s-cis* conformation with the O2—C7—C8—C9 torsion angle being 6.5 (5) $^{\circ}$  for molecule A and 8.8 (5) $^{\circ}$  for B. In what follows, the distortion of the angles is relative to what is expected in terms of hybridization principles. In molecule A, the slight enlargement of the C5A—C6A—C7A (122.4 (3) $^{\circ}$ ) and C6A—C7A—C8A (121.9 (3) $^{\circ}$ ) angles may be the result of the short H5AA···H8AA (2.14 Å) contact whereas the short H8AA···H11A (2.35 Å) contact may widen the C8A—C9A—C10A (129.0 (3) $^{\circ}$ ) and C9A—C10A—C11A (123.0 (3) $^{\circ}$ ) angles. The short H14A···H16A (2.26 Å) contact may result in the opening of the O3A—C13A—C14A (124.9 (3) $^{\circ}$ ) angle. Likewise, in molecule B, a close interatomic contact between H5BA and H8BA (2.13 Å) may result in the widening of the C5B—C6B—C7B (123.1 (3) $^{\circ}$ ) and C6B—C7B—C8B (121.2 (3) $^{\circ}$ ) angles whereas the opening of C8B—C9B—C10B and C9B—C10B—C11B angles to 128.6 (3) $^{\circ}$  and 123.0 (3) $^{\circ}$ , respectively, may be the result of the close H8BA···H11B (2.30 Å) contact. Similar strain induced by a short H14B···H16C (2.26 Å) contact may result in the opening of the O3B—C13B—C14B (125.2 (3) $^{\circ}$ ) angle. These features were also reported in related structures (Ng *et al.*, 2006; Razak *et al.*, 2009; Ngaini, Fadzillah *et al.*, 2009; Ngaini, Rahman *et al.*, 2009).

The conformation throughout the zigzag alkoxy tails in both molecules is *trans* with the largest deviation from the ideal value being -174.4 (3) $^{\circ}$  for the C19A—C20A—C21A—C22A torsion angle in molecule A and -173.5 (3) $^{\circ}$  for the C17B—C18B—C19B—C20B torsion angle in B. Even though the torsion angle C16—O3—C13—C14 in each molecule is -6.8 (5) $^{\circ}$  for A and -12.2 (5) $^{\circ}$  for B, the alkoxy chains curve out of the zigzag plane with the least-squares plane through the chain making dihedral angle with the attached benzene ring of 17.02 (19) $^{\circ}$  [maximum deviation of -0.302 (4) Å at C21A] and 16.73 (19) $^{\circ}$  [maximum deviation of -0.256 (4) Å at C21B], for molecules A and B, respectively.

An intramolecular O-H···O interaction involving the keto and hydroxy groups (Table 1) in both molecules generates S(6) ring motifs (Bernstein *et al.*, 1995). In the crystal structure, the molecules are stacked alternately along the *a* axis in a

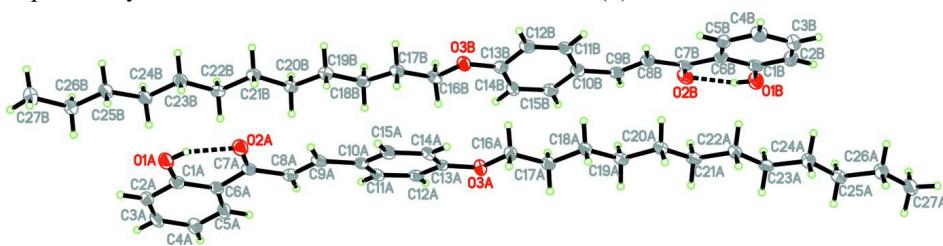
head-to-tail manner (Fig. 2). In the absence of conventional hydrogen bonds, the crystal structure is stabilized by weak C—H···π interactions (Table 1).

## S2. Experimental

A mixture of 2-hydroxyacetophenone (2.72 ml, 20 mmol) and 4-dodecyloxybenzaldehyde (5.81 ml, 20 mmol) and KOH (4.04 g, 72 mmol) in 60 ml of methanol was heated at reflux for 10 h. The reaction was cooled to room temperature and acidified with cold diluted HCl (2 N). The resulting precipitate was filtered, washed and dried. After redissolving in hexane, followed by a few days of slow evaporation, crystals were collected.

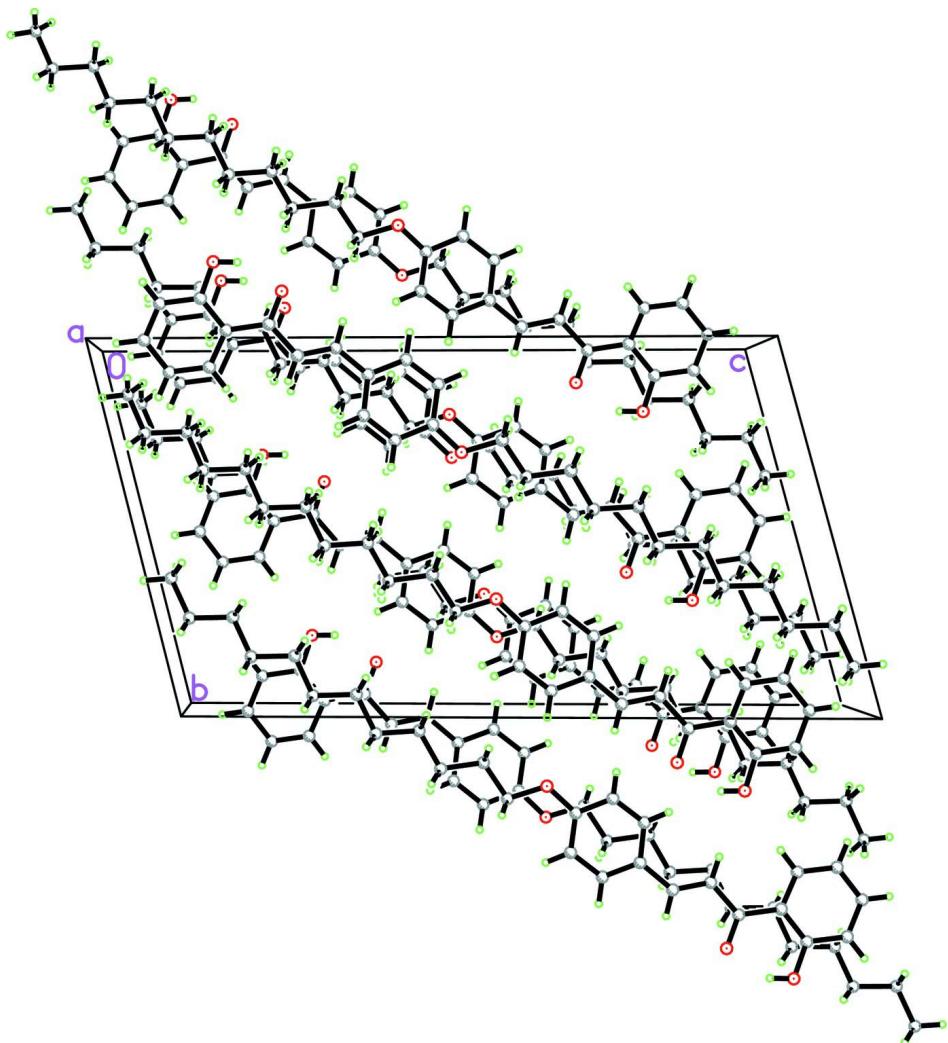
## S3. Refinement

All H atoms were positioned geometrically and refined using a riding model. The  $U_{\text{iso}}(\text{H})$  values were constrained to be  $1.5U_{\text{eq}}(\text{C},\text{O})$  (methyl H and hydroxyl H atoms) and  $1.2U_{\text{eq}}(\text{C})$  (other H atoms). The rotating model group was considered for the methyl group. The crystal is a twin with a refined BASF = 0.212 (2).



**Figure 1**

The asymmetric unit of the title compound, showing 50% probability displacement ellipsoids and the atom numbering scheme. Intramolecular interactions were shown as dashed line.

**Figure 2**

Part of the crystal structure of (I) viewed along the *a* axis.

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#### Crystal data

$C_{27}H_{36}O_3$   
 $M_r = 408.56$   
Triclinic,  $P\bar{1}$   
Hall symbol: -P 1  
 $a = 7.4953 (6)$  Å  
 $b = 13.4714 (9)$  Å  
 $c = 23.7874 (18)$  Å  
 $\alpha = 75.116 (4)^\circ$   
 $\beta = 83.876 (5)^\circ$   
 $\gamma = 84.669 (5)^\circ$   
 $V = 2302.7 (3)$  Å<sup>3</sup>

$Z = 4$   
 $F(000) = 888$   
 $D_x = 1.179$  Mg m<sup>-3</sup>  
Mo  $K\alpha$  radiation,  $\lambda = 0.71073$  Å  
Cell parameters from 3668 reflections  
 $\theta = 2.7\text{--}28.0^\circ$   
 $\mu = 0.08$  mm<sup>-1</sup>  
 $T = 100$  K  
Needle, yellow  
 $0.55 \times 0.13 \times 0.06$  mm

*Data collection*

Bruker APEXII  
diffractometer  
Radiation source: sealed tube  
Graphite monochromator  
 $\pi$  and  $\omega$  scans  
Absorption correction: multi-scan  
(SADABS; Bruker, 2005)  
 $T_{\min} = 0.960$ ,  $T_{\max} = 0.996$

8571 measured reflections  
8571 independent reflections  
4737 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.000$   
 $\theta_{\max} = 25.5^\circ$ ,  $\theta_{\min} = 0.9^\circ$   
 $h = -8 \rightarrow 9$   
 $k = -16 \rightarrow 16$   
 $l = -28 \rightarrow 28$

*Refinement*

Refinement on  $F^2$   
Least-squares matrix: full  
 $R[F^2 > 2\sigma(F^2)] = 0.069$   
 $wR(F^2) = 0.206$   
 $S = 1.03$   
8571 reflections  
546 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  
 $w = 1/[\sigma^2(F_o^2) + (0.0924P)^2 + 0.763P]$   
where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} < 0.001$   
 $\Delta\rho_{\max} = 0.29 \text{ e } \text{\AA}^{-3}$   
 $\Delta\rho_{\min} = -0.30 \text{ e } \text{\AA}^{-3}$

*Special details*

**Experimental.** The crystal was placed in the cold stream of an Oxford Cryosystems Cobra open-flow nitrogen cryostat (Cosier & Glazer, 1986) operating at 100.0 (1) K.

**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 ( $\text{\AA}^2$ )*

|      | $x$        | $y$          | $z$          | $U_{\text{iso}}^*/U_{\text{eq}}$ |
|------|------------|--------------|--------------|----------------------------------|
| O1A  | 0.6765 (4) | 0.69790 (18) | 0.78918 (11) | 0.0307 (6)                       |
| H1A  | 0.6609     | 0.6985       | 0.7555       | 0.046*                           |
| O2A  | 0.6335 (3) | 0.62359 (18) | 0.70527 (11) | 0.0290 (6)                       |
| O3A  | 0.6291 (3) | 0.19687 (18) | 0.50595 (10) | 0.0255 (6)                       |
| C1A  | 0.6917 (5) | 0.5994 (3)   | 0.82144 (16) | 0.0249 (8)                       |
| C2A  | 0.7137 (5) | 0.5837 (3)   | 0.88068 (16) | 0.0273 (9)                       |
| H2AA | 0.7158     | 0.6397       | 0.8966       | 0.033*                           |
| C3A  | 0.7324 (5) | 0.4852 (3)   | 0.91544 (16) | 0.0294 (9)                       |
| H3AA | 0.7456     | 0.4751       | 0.9550       | 0.035*                           |
| C4A  | 0.7320 (5) | 0.4007 (3)   | 0.89221 (15) | 0.0288 (9)                       |
| H4AA | 0.7469     | 0.3344       | 0.9159       | 0.035*                           |
| C5A  | 0.7094 (5) | 0.4154 (3)   | 0.83407 (15) | 0.0256 (8)                       |
| H5AA | 0.7088     | 0.3585       | 0.8187       | 0.031*                           |
| C6A  | 0.6871 (4) | 0.5145 (3)   | 0.79750 (15) | 0.0210 (8)                       |

|      |            |             |              |            |
|------|------------|-------------|--------------|------------|
| C7A  | 0.6563 (4) | 0.5329 (3)  | 0.73494 (15) | 0.0220 (8) |
| C8A  | 0.6552 (5) | 0.4480 (3)  | 0.70683 (15) | 0.0235 (8) |
| H8AA | 0.6597     | 0.3802      | 0.7291       | 0.028*     |
| C9A  | 0.6477 (4) | 0.4687 (3)  | 0.64905 (15) | 0.0231 (8) |
| H9AA | 0.6432     | 0.5381      | 0.6297       | 0.028*     |
| C10A | 0.6455 (4) | 0.3974 (3)  | 0.61220 (15) | 0.0214 (8) |
| C11A | 0.6156 (5) | 0.2928 (3)  | 0.63470 (15) | 0.0230 (8) |
| H11A | 0.5988     | 0.2660      | 0.6749       | 0.028*     |
| C12A | 0.6109 (4) | 0.2297 (3)  | 0.59822 (15) | 0.0221 (8) |
| H12A | 0.5905     | 0.1606      | 0.6139       | 0.027*     |
| C13A | 0.6362 (4) | 0.2674 (3)  | 0.53777 (15) | 0.0203 (8) |
| C14A | 0.6673 (5) | 0.3704 (3)  | 0.51432 (15) | 0.0229 (8) |
| H14A | 0.6862     | 0.3961      | 0.4741       | 0.028*     |
| C15A | 0.6698 (5) | 0.4344 (3)  | 0.55132 (15) | 0.0233 (8) |
| H15A | 0.6882     | 0.5037      | 0.5354       | 0.028*     |
| C16A | 0.6352 (5) | 0.2302 (3)  | 0.44325 (15) | 0.0236 (8) |
| H16A | 0.7410     | 0.2680      | 0.4277       | 0.028*     |
| H16B | 0.5292     | 0.2746      | 0.4313       | 0.028*     |
| C17A | 0.6417 (5) | 0.1341 (3)  | 0.42151 (15) | 0.0252 (8) |
| H17A | 0.5401     | 0.0951      | 0.4406       | 0.030*     |
| H17B | 0.7502     | 0.0923      | 0.4334       | 0.030*     |
| C18A | 0.6386 (5) | 0.1513 (3)  | 0.35641 (15) | 0.0233 (8) |
| H18A | 0.7441     | 0.1861      | 0.3368       | 0.028*     |
| H18B | 0.5331     | 0.1953      | 0.3437       | 0.028*     |
| C19A | 0.6354 (5) | 0.0500 (3)  | 0.33918 (15) | 0.0240 (8) |
| H19A | 0.7360     | 0.0046      | 0.3550       | 0.029*     |
| H19B | 0.5261     | 0.0178      | 0.3575       | 0.029*     |
| C20A | 0.6443 (5) | 0.0575 (3)  | 0.27412 (15) | 0.0255 (8) |
| H20A | 0.7542     | 0.0885      | 0.2554       | 0.031*     |
| H20B | 0.5437     | 0.1023      | 0.2578       | 0.031*     |
| C21A | 0.6396 (5) | -0.0469 (3) | 0.26067 (15) | 0.0252 (8) |
| H21A | 0.7325     | -0.0935     | 0.2806       | 0.030*     |
| H21B | 0.5246     | -0.0746     | 0.2764       | 0.030*     |
| C22A | 0.6664 (5) | -0.0442 (3) | 0.19601 (15) | 0.0246 (8) |
| H22A | 0.7767     | -0.0116     | 0.1794       | 0.029*     |
| H22B | 0.5678     | -0.0025     | 0.1765       | 0.029*     |
| C23A | 0.6765 (5) | -0.1509 (3) | 0.18409 (15) | 0.0251 (8) |
| H23A | 0.7717     | -0.1936     | 0.2051       | 0.030*     |
| H23B | 0.5641     | -0.1823     | 0.1992       | 0.030*     |
| C24A | 0.7112 (5) | -0.1494 (3) | 0.11987 (15) | 0.0265 (9) |
| H24A | 0.8193     | -0.1139     | 0.1041       | 0.032*     |
| H24B | 0.6119     | -0.1105     | 0.0993       | 0.032*     |
| C25A | 0.7332 (5) | -0.2558 (3) | 0.10808 (15) | 0.0276 (9) |
| H25A | 0.8331     | -0.2946     | 0.1284       | 0.033*     |
| H25B | 0.6254     | -0.2915     | 0.1240       | 0.033*     |
| C26A | 0.7664 (5) | -0.2538 (3) | 0.04410 (15) | 0.0288 (9) |
| H26A | 0.8748     | -0.2187     | 0.0282       | 0.035*     |
| H26B | 0.6671     | -0.2145     | 0.0237       | 0.035*     |

|      |            |               |              |             |
|------|------------|---------------|--------------|-------------|
| C27A | 0.7868 (6) | -0.3608 (3)   | 0.03238 (16) | 0.0336 (10) |
| H27A | 0.8210     | -0.3548       | -0.0084      | 0.050*      |
| H27B | 0.6745     | -0.3925       | 0.0431       | 0.050*      |
| H27C | 0.8778     | -0.4023       | 0.0551       | 0.050*      |
| O1B  | 0.1478 (4) | -0.19486 (18) | 0.21550 (11) | 0.0324 (7)  |
| H1B  | 0.1245     | -0.1949       | 0.2500       | 0.049*      |
| O2B  | 0.0945 (3) | -0.11900 (18) | 0.30324 (11) | 0.0290 (6)  |
| O3B  | 0.1502 (3) | 0.30711 (17)  | 0.50023 (10) | 0.0251 (6)  |
| C1B  | 0.1699 (5) | -0.0980 (3)   | 0.18370 (16) | 0.0248 (8)  |
| C2B  | 0.2029 (5) | -0.0831 (3)   | 0.12353 (17) | 0.0296 (9)  |
| H2BA | 0.2081     | -0.1390       | 0.1071       | 0.036*      |
| C3B  | 0.2280 (5) | 0.0137 (3)    | 0.08821 (17) | 0.0328 (10) |
| H3BA | 0.2487     | 0.0230        | 0.0480       | 0.039*      |
| C4B  | 0.2224 (5) | 0.0976 (3)    | 0.11249 (16) | 0.0310 (9)  |
| H4BA | 0.2395     | 0.1631        | 0.0886       | 0.037*      |
| C5B  | 0.1917 (5) | 0.0837 (3)    | 0.17155 (16) | 0.0252 (8)  |
| H5BA | 0.1892     | 0.1402        | 0.1874       | 0.030*      |
| C6B  | 0.1638 (4) | -0.0137 (3)   | 0.20895 (16) | 0.0224 (8)  |
| C7B  | 0.1286 (4) | -0.0313 (3)   | 0.27275 (15) | 0.0224 (8)  |
| C8B  | 0.1362 (4) | 0.0522 (3)    | 0.30173 (15) | 0.0230 (8)  |
| H8BA | 0.1448     | 0.1195        | 0.2794       | 0.028*      |
| C9B  | 0.1308 (4) | 0.0320 (3)    | 0.35977 (15) | 0.0217 (8)  |
| H9BA | 0.1227     | -0.0367       | 0.3796       | 0.026*      |
| C10B | 0.1359 (5) | 0.1032 (3)    | 0.39596 (15) | 0.0231 (8)  |
| C11B | 0.1184 (5) | 0.2102 (3)    | 0.37436 (15) | 0.0234 (8)  |
| H11B | 0.1037     | 0.2379        | 0.3350       | 0.028*      |
| C12B | 0.1225 (5) | 0.2748 (3)    | 0.41016 (16) | 0.0258 (9)  |
| H12B | 0.1100     | 0.3457        | 0.3949       | 0.031*      |
| C13B | 0.1452 (5) | 0.2351 (3)    | 0.46948 (16) | 0.0230 (8)  |
| C14B | 0.1622 (5) | 0.1297 (3)    | 0.49237 (16) | 0.0249 (8)  |
| H14B | 0.1760     | 0.1024        | 0.5319       | 0.030*      |
| C15B | 0.1585 (5) | 0.0651 (3)    | 0.45546 (16) | 0.0259 (9)  |
| H15B | 0.1713     | -0.0057       | 0.4708       | 0.031*      |
| C16B | 0.1390 (5) | 0.2755 (3)    | 0.56305 (15) | 0.0237 (8)  |
| H16C | 0.2351     | 0.2240        | 0.5761       | 0.028*      |
| H16D | 0.0247     | 0.2466        | 0.5783       | 0.028*      |
| C17B | 0.1564 (5) | 0.3710 (3)    | 0.58335 (15) | 0.0247 (8)  |
| H17C | 0.0658     | 0.4232        | 0.5666       | 0.030*      |
| H17D | 0.2732     | 0.3969        | 0.5684       | 0.030*      |
| C18B | 0.1366 (5) | 0.3554 (3)    | 0.64911 (15) | 0.0245 (8)  |
| H18C | 0.2292     | 0.3050        | 0.6663       | 0.029*      |
| H18D | 0.0206     | 0.3289        | 0.6646       | 0.029*      |
| C19B | 0.1520 (5) | 0.4559 (3)    | 0.66580 (15) | 0.0249 (8)  |
| H19C | 0.2613     | 0.4858        | 0.6457       | 0.030*      |
| H19D | 0.0517     | 0.5031        | 0.6515       | 0.030*      |
| C20B | 0.1549 (5) | 0.4484 (3)    | 0.73017 (15) | 0.0231 (8)  |
| H20C | 0.2600     | 0.4055        | 0.7444       | 0.028*      |
| H20D | 0.0492     | 0.4153        | 0.7510       | 0.028*      |

|      |            |            |              |             |
|------|------------|------------|--------------|-------------|
| C21B | 0.1587 (5) | 0.5525 (3) | 0.74315 (15) | 0.0248 (8)  |
| H21C | 0.0480     | 0.5927     | 0.7321       | 0.030*      |
| H21D | 0.2572     | 0.5882     | 0.7189       | 0.030*      |
| C22B | 0.1799 (5) | 0.5491 (3) | 0.80653 (15) | 0.0255 (8)  |
| H22C | 0.0780     | 0.5170     | 0.8307       | 0.031*      |
| H22D | 0.2875     | 0.5063     | 0.8184       | 0.031*      |
| C23B | 0.1928 (5) | 0.6542 (3) | 0.81750 (15) | 0.0262 (8)  |
| H23C | 0.0837     | 0.6963     | 0.8067       | 0.031*      |
| H23D | 0.2924     | 0.6871     | 0.7924       | 0.031*      |
| C24B | 0.2195 (5) | 0.6512 (3) | 0.88035 (15) | 0.0262 (8)  |
| H24C | 0.3210     | 0.6032     | 0.8925       | 0.031*      |
| H24D | 0.1137     | 0.6252     | 0.9049       | 0.031*      |
| C25B | 0.2525 (5) | 0.7555 (3) | 0.89035 (15) | 0.0276 (9)  |
| H25C | 0.3601     | 0.7808     | 0.8666       | 0.033*      |
| H25D | 0.1524     | 0.8041     | 0.8774       | 0.033*      |
| C26B | 0.2743 (5) | 0.7519 (3) | 0.95361 (15) | 0.0281 (9)  |
| H26C | 0.3818     | 0.7090     | 0.9654       | 0.034*      |
| H26D | 0.1724     | 0.7200     | 0.9779       | 0.034*      |
| C27B | 0.2879 (6) | 0.8576 (3) | 0.96430 (17) | 0.0361 (10) |
| H27D | 0.3122     | 0.8498     | 1.0041       | 0.054*      |
| H27E | 0.1764     | 0.8979     | 0.9570       | 0.054*      |
| H27F | 0.3835     | 0.8915     | 0.9386       | 0.054*      |

*Atomic displacement parameters ( $\text{\AA}^2$ )*

|      | $U^{11}$    | $U^{22}$    | $U^{33}$    | $U^{12}$     | $U^{13}$     | $U^{23}$     |
|------|-------------|-------------|-------------|--------------|--------------|--------------|
| O1A  | 0.0346 (16) | 0.0267 (14) | 0.0343 (16) | -0.0020 (11) | -0.0117 (13) | -0.0102 (12) |
| O2A  | 0.0320 (15) | 0.0243 (14) | 0.0325 (15) | -0.0041 (11) | -0.0084 (12) | -0.0074 (11) |
| O3A  | 0.0314 (15) | 0.0256 (14) | 0.0220 (14) | -0.0052 (11) | -0.0045 (11) | -0.0081 (11) |
| C1A  | 0.0151 (19) | 0.027 (2)   | 0.033 (2)   | -0.0025 (15) | -0.0031 (16) | -0.0072 (17) |
| C2A  | 0.022 (2)   | 0.031 (2)   | 0.034 (2)   | -0.0024 (16) | -0.0058 (17) | -0.0160 (17) |
| C3A  | 0.025 (2)   | 0.043 (2)   | 0.023 (2)   | -0.0046 (17) | -0.0076 (17) | -0.0121 (18) |
| C4A  | 0.032 (2)   | 0.029 (2)   | 0.025 (2)   | -0.0052 (17) | -0.0047 (17) | -0.0054 (16) |
| C5A  | 0.025 (2)   | 0.027 (2)   | 0.027 (2)   | -0.0053 (16) | -0.0017 (16) | -0.0102 (16) |
| C6A  | 0.0143 (18) | 0.027 (2)   | 0.024 (2)   | -0.0053 (15) | -0.0010 (15) | -0.0078 (16) |
| C7A  | 0.0136 (18) | 0.025 (2)   | 0.027 (2)   | -0.0029 (14) | -0.0030 (15) | -0.0055 (16) |
| C8A  | 0.020 (2)   | 0.024 (2)   | 0.027 (2)   | -0.0006 (15) | -0.0037 (16) | -0.0048 (16) |
| C9A  | 0.0148 (19) | 0.025 (2)   | 0.029 (2)   | -0.0038 (14) | -0.0057 (16) | -0.0032 (16) |
| C10A | 0.0111 (18) | 0.028 (2)   | 0.028 (2)   | -0.0022 (14) | -0.0037 (15) | -0.0102 (16) |
| C11A | 0.0179 (19) | 0.029 (2)   | 0.023 (2)   | -0.0024 (15) | -0.0054 (15) | -0.0049 (16) |
| C12A | 0.0167 (19) | 0.0210 (19) | 0.028 (2)   | -0.0016 (14) | -0.0059 (15) | -0.0042 (16) |
| C13A | 0.0132 (18) | 0.0224 (19) | 0.029 (2)   | 0.0012 (14)  | -0.0062 (15) | -0.0130 (16) |
| C14A | 0.022 (2)   | 0.027 (2)   | 0.021 (2)   | -0.0014 (15) | -0.0082 (16) | -0.0062 (16) |
| C15A | 0.0210 (19) | 0.0197 (19) | 0.029 (2)   | -0.0052 (15) | -0.0059 (16) | -0.0035 (15) |
| C16A | 0.022 (2)   | 0.026 (2)   | 0.025 (2)   | -0.0033 (15) | -0.0034 (16) | -0.0098 (16) |
| C17A | 0.022 (2)   | 0.030 (2)   | 0.025 (2)   | -0.0019 (16) | -0.0032 (16) | -0.0084 (16) |
| C18A | 0.021 (2)   | 0.025 (2)   | 0.026 (2)   | -0.0010 (15) | -0.0059 (16) | -0.0073 (16) |
| C19A | 0.021 (2)   | 0.026 (2)   | 0.025 (2)   | -0.0035 (15) | -0.0063 (16) | -0.0045 (16) |

|      |             |             |             |              |              |              |
|------|-------------|-------------|-------------|--------------|--------------|--------------|
| C20A | 0.020 (2)   | 0.030 (2)   | 0.028 (2)   | -0.0011 (16) | -0.0066 (16) | -0.0077 (16) |
| C21A | 0.024 (2)   | 0.0225 (19) | 0.031 (2)   | -0.0004 (15) | -0.0100 (17) | -0.0057 (16) |
| C22A | 0.022 (2)   | 0.028 (2)   | 0.026 (2)   | -0.0028 (15) | -0.0058 (16) | -0.0092 (16) |
| C23A | 0.021 (2)   | 0.027 (2)   | 0.027 (2)   | 0.0012 (15)  | -0.0087 (16) | -0.0061 (16) |
| C24A | 0.026 (2)   | 0.029 (2)   | 0.025 (2)   | -0.0020 (16) | -0.0028 (16) | -0.0075 (16) |
| C25A | 0.026 (2)   | 0.030 (2)   | 0.028 (2)   | -0.0037 (16) | -0.0076 (17) | -0.0067 (16) |
| C26A | 0.032 (2)   | 0.029 (2)   | 0.026 (2)   | -0.0017 (17) | -0.0053 (17) | -0.0075 (16) |
| C27A | 0.040 (3)   | 0.036 (2)   | 0.028 (2)   | -0.0003 (18) | -0.0104 (19) | -0.0130 (18) |
| O1B  | 0.0386 (17) | 0.0279 (15) | 0.0329 (16) | -0.0060 (12) | -0.0052 (14) | -0.0094 (12) |
| O2B  | 0.0291 (15) | 0.0265 (14) | 0.0324 (15) | -0.0084 (11) | -0.0100 (12) | -0.0040 (12) |
| O3B  | 0.0280 (15) | 0.0236 (13) | 0.0250 (14) | -0.0037 (11) | -0.0058 (11) | -0.0064 (11) |
| C1B  | 0.0171 (19) | 0.027 (2)   | 0.032 (2)   | -0.0024 (15) | -0.0081 (16) | -0.0070 (17) |
| C2B  | 0.029 (2)   | 0.029 (2)   | 0.036 (2)   | -0.0018 (17) | -0.0062 (18) | -0.0156 (18) |
| C3B  | 0.034 (2)   | 0.040 (2)   | 0.026 (2)   | 0.0029 (18)  | -0.0060 (18) | -0.0129 (18) |
| C4B  | 0.034 (2)   | 0.027 (2)   | 0.030 (2)   | -0.0030 (17) | -0.0050 (18) | -0.0033 (17) |
| C5B  | 0.025 (2)   | 0.023 (2)   | 0.030 (2)   | 0.0008 (15)  | -0.0060 (17) | -0.0105 (16) |
| C6B  | 0.0127 (18) | 0.0245 (19) | 0.032 (2)   | 0.0008 (14)  | -0.0059 (16) | -0.0104 (16) |
| C7B  | 0.0121 (18) | 0.027 (2)   | 0.029 (2)   | -0.0004 (14) | -0.0098 (15) | -0.0060 (16) |
| C8B  | 0.0172 (19) | 0.0221 (19) | 0.031 (2)   | -0.0010 (15) | -0.0056 (16) | -0.0068 (16) |
| C9B  | 0.0161 (19) | 0.0212 (19) | 0.028 (2)   | -0.0028 (14) | -0.0046 (15) | -0.0053 (15) |
| C10B | 0.0159 (19) | 0.025 (2)   | 0.030 (2)   | -0.0054 (15) | -0.0028 (16) | -0.0078 (16) |
| C11B | 0.0180 (19) | 0.030 (2)   | 0.022 (2)   | -0.0036 (15) | -0.0038 (15) | -0.0049 (16) |
| C12B | 0.019 (2)   | 0.026 (2)   | 0.033 (2)   | -0.0031 (15) | -0.0057 (16) | -0.0059 (17) |
| C13B | 0.0152 (19) | 0.027 (2)   | 0.028 (2)   | -0.0052 (15) | -0.0021 (15) | -0.0082 (16) |
| C14B | 0.025 (2)   | 0.029 (2)   | 0.021 (2)   | -0.0026 (16) | -0.0044 (16) | -0.0067 (16) |
| C15B | 0.023 (2)   | 0.024 (2)   | 0.030 (2)   | -0.0020 (15) | -0.0055 (17) | -0.0046 (16) |
| C16B | 0.0204 (19) | 0.026 (2)   | 0.024 (2)   | -0.0050 (15) | -0.0049 (15) | -0.0031 (16) |
| C17B | 0.0174 (19) | 0.028 (2)   | 0.029 (2)   | -0.0025 (15) | -0.0041 (16) | -0.0058 (16) |
| C18B | 0.020 (2)   | 0.027 (2)   | 0.026 (2)   | -0.0006 (15) | -0.0043 (16) | -0.0061 (16) |
| C19B | 0.022 (2)   | 0.025 (2)   | 0.027 (2)   | -0.0014 (15) | -0.0054 (16) | -0.0034 (16) |
| C20B | 0.0196 (19) | 0.025 (2)   | 0.024 (2)   | -0.0023 (15) | -0.0044 (15) | -0.0046 (15) |
| C21B | 0.019 (2)   | 0.027 (2)   | 0.028 (2)   | -0.0025 (15) | -0.0023 (16) | -0.0061 (16) |
| C22B | 0.021 (2)   | 0.028 (2)   | 0.028 (2)   | 0.0007 (15)  | -0.0052 (16) | -0.0069 (16) |
| C23B | 0.024 (2)   | 0.025 (2)   | 0.030 (2)   | -0.0040 (16) | -0.0052 (17) | -0.0072 (16) |
| C24B | 0.025 (2)   | 0.027 (2)   | 0.028 (2)   | -0.0034 (16) | -0.0079 (17) | -0.0068 (16) |
| C25B | 0.027 (2)   | 0.029 (2)   | 0.028 (2)   | -0.0027 (16) | -0.0070 (17) | -0.0079 (16) |
| C26B | 0.030 (2)   | 0.031 (2)   | 0.024 (2)   | -0.0031 (17) | -0.0047 (17) | -0.0084 (16) |
| C27B | 0.044 (3)   | 0.035 (2)   | 0.033 (2)   | -0.0091 (19) | -0.006 (2)   | -0.0123 (18) |

Geometric parameters ( $\text{\AA}$ ,  $^\circ$ )

|          |           |          |           |
|----------|-----------|----------|-----------|
| O1A—C1A  | 1.353 (4) | O1B—C1B  | 1.345 (4) |
| O1A—H1A  | 0.8200    | O1B—H1B  | 0.8200    |
| O2A—C7A  | 1.251 (4) | O2B—C7B  | 1.251 (4) |
| O3A—C13A | 1.366 (4) | O3B—C13B | 1.361 (4) |
| O3A—C16A | 1.440 (4) | O3B—C16B | 1.440 (4) |
| C1A—C2A  | 1.396 (5) | C1B—C2B  | 1.392 (5) |
| C1A—C6A  | 1.407 (5) | C1B—C6B  | 1.410 (5) |

|           |           |           |           |
|-----------|-----------|-----------|-----------|
| C2A—C3A   | 1.374 (5) | C2B—C3B   | 1.375 (5) |
| C2A—H2AA  | 0.9300    | C2B—H2BA  | 0.9300    |
| C3A—C4A   | 1.388 (5) | C3B—C4B   | 1.392 (5) |
| C3A—H3AA  | 0.9300    | C3B—H3BA  | 0.9300    |
| C4A—C5A   | 1.372 (5) | C4B—C5B   | 1.366 (5) |
| C4A—H4AA  | 0.9300    | C4B—H4BA  | 0.9300    |
| C5A—C6A   | 1.400 (5) | C5B—C6B   | 1.403 (5) |
| C5A—H5AA  | 0.9300    | C5B—H5BA  | 0.9300    |
| C6A—C7A   | 1.484 (5) | C6B—C7B   | 1.473 (5) |
| C7A—C8A   | 1.467 (5) | C7B—C8B   | 1.470 (5) |
| C8A—C9A   | 1.337 (5) | C8B—C9B   | 1.334 (5) |
| C8A—H8AA  | 0.9300    | C8B—H8BA  | 0.9300    |
| C9A—C10A  | 1.459 (5) | C9B—C10B  | 1.449 (5) |
| C9A—H9AA  | 0.9300    | C9B—H9BA  | 0.9300    |
| C10A—C15A | 1.403 (5) | C10B—C15B | 1.399 (5) |
| C10A—C11A | 1.404 (5) | C10B—C11B | 1.399 (5) |
| C11A—C12A | 1.366 (5) | C11B—C12B | 1.370 (5) |
| C11A—H11A | 0.9300    | C11B—H11B | 0.9300    |
| C12A—C13A | 1.395 (5) | C12B—C13B | 1.397 (5) |
| C12A—H12A | 0.9300    | C12B—H12B | 0.9300    |
| C13A—C14A | 1.387 (5) | C13B—C14B | 1.384 (5) |
| C14A—C15A | 1.384 (5) | C14B—C15B | 1.389 (5) |
| C14A—H14A | 0.9300    | C14B—H14B | 0.9300    |
| C15A—H15A | 0.9300    | C15B—H15B | 0.9300    |
| C16A—C17A | 1.508 (5) | C16B—C17B | 1.507 (5) |
| C16A—H16A | 0.9700    | C16B—H16C | 0.9700    |
| C16A—H16B | 0.9700    | C16B—H16D | 0.9700    |
| C17A—C18A | 1.508 (5) | C17B—C18B | 1.517 (5) |
| C17A—H17A | 0.9700    | C17B—H17C | 0.9700    |
| C17A—H17B | 0.9700    | C17B—H17D | 0.9700    |
| C18A—C19A | 1.525 (5) | C18B—C19B | 1.522 (5) |
| C18A—H18A | 0.9700    | C18B—H18C | 0.9700    |
| C18A—H18B | 0.9700    | C18B—H18D | 0.9700    |
| C19A—C20A | 1.519 (5) | C19B—C20B | 1.511 (5) |
| C19A—H19A | 0.9700    | C19B—H19C | 0.9700    |
| C19A—H19B | 0.9700    | C19B—H19D | 0.9700    |
| C20A—C21A | 1.525 (5) | C20B—C21B | 1.515 (5) |
| C20A—H20A | 0.9700    | C20B—H20C | 0.9700    |
| C20A—H20B | 0.9700    | C20B—H20D | 0.9700    |
| C21A—C22A | 1.522 (5) | C21B—C22B | 1.521 (5) |
| C21A—H21A | 0.9700    | C21B—H21C | 0.9700    |
| C21A—H21B | 0.9700    | C21B—H21D | 0.9700    |
| C22A—C23A | 1.529 (5) | C22B—C23B | 1.518 (5) |
| C22A—H22A | 0.9700    | C22B—H22C | 0.9700    |
| C22A—H22B | 0.9700    | C22B—H22D | 0.9700    |
| C23A—C24A | 1.517 (5) | C23B—C24B | 1.519 (5) |
| C23A—H23A | 0.9700    | C23B—H23C | 0.9700    |
| C23A—H23B | 0.9700    | C23B—H23D | 0.9700    |

|                |           |                |           |
|----------------|-----------|----------------|-----------|
| C24A—C25A      | 1.520 (5) | C24B—C25B      | 1.531 (5) |
| C24A—H24A      | 0.9700    | C24B—H24C      | 0.9700    |
| C24A—H24B      | 0.9700    | C24B—H24D      | 0.9700    |
| C25A—C26A      | 1.510 (5) | C25B—C26B      | 1.519 (5) |
| C25A—H25A      | 0.9700    | C25B—H25C      | 0.9700    |
| C25A—H25B      | 0.9700    | C25B—H25D      | 0.9700    |
| C26A—C27A      | 1.528 (5) | C26B—C27B      | 1.523 (5) |
| C26A—H26A      | 0.9700    | C26B—H26C      | 0.9700    |
| C26A—H26B      | 0.9700    | C26B—H26D      | 0.9700    |
| C27A—H27A      | 0.9600    | C27B—H27D      | 0.9600    |
| C27A—H27B      | 0.9600    | C27B—H27E      | 0.9600    |
| C27A—H27C      | 0.9600    | C27B—H27F      | 0.9600    |
| <br>           |           |                |           |
| C1A—O1A—H1A    | 109.5     | C1B—O1B—H1B    | 109.5     |
| C13A—O3A—C16A  | 119.7 (3) | C13B—O3B—C16B  | 119.3 (3) |
| O1A—C1A—C2A    | 117.3 (3) | O1B—C1B—C2B    | 117.2 (3) |
| O1A—C1A—C6A    | 122.7 (3) | O1B—C1B—C6B    | 122.8 (3) |
| C2A—C1A—C6A    | 120.0 (3) | C2B—C1B—C6B    | 120.0 (3) |
| C3A—C2A—C1A    | 119.9 (3) | C3B—C2B—C1B    | 120.5 (3) |
| C3A—C2A—H2AA   | 120.0     | C3B—C2B—H2BA   | 119.7     |
| C1A—C2A—H2AA   | 120.0     | C1B—C2B—H2BA   | 119.7     |
| C2A—C3A—C4A    | 120.8 (3) | C2B—C3B—C4B    | 120.1 (4) |
| C2A—C3A—H3AA   | 119.6     | C2B—C3B—H3BA   | 119.9     |
| C4A—C3A—H3AA   | 119.6     | C4B—C3B—H3BA   | 119.9     |
| C5A—C4A—C3A    | 119.7 (3) | C5B—C4B—C3B    | 119.8 (3) |
| C5A—C4A—H4AA   | 120.1     | C5B—C4B—H4BA   | 120.1     |
| C3A—C4A—H4AA   | 120.1     | C3B—C4B—H4BA   | 120.1     |
| C4A—C5A—C6A    | 121.1 (3) | C4B—C5B—C6B    | 121.7 (3) |
| C4A—C5A—H5AA   | 119.5     | C4B—C5B—H5BA   | 119.2     |
| C6A—C5A—H5AA   | 119.5     | C6B—C5B—H5BA   | 119.2     |
| C5A—C6A—C1A    | 118.5 (3) | C5B—C6B—C1B    | 117.8 (3) |
| C5A—C6A—C7A    | 122.4 (3) | C5B—C6B—C7B    | 123.1 (3) |
| C1A—C6A—C7A    | 119.1 (3) | C1B—C6B—C7B    | 119.1 (3) |
| O2A—C7A—C8A    | 119.3 (3) | O2B—C7B—C8B    | 119.0 (3) |
| O2A—C7A—C6A    | 118.8 (3) | O2B—C7B—C6B    | 119.8 (3) |
| C8A—C7A—C6A    | 121.9 (3) | C8B—C7B—C6B    | 121.2 (3) |
| C9A—C8A—C7A    | 119.6 (3) | C9B—C8B—C7B    | 120.5 (3) |
| C9A—C8A—H8AA   | 120.2     | C9B—C8B—H8BA   | 119.7     |
| C7A—C8A—H8AA   | 120.2     | C7B—C8B—H8BA   | 119.7     |
| C8A—C9A—C10A   | 129.0 (3) | C8B—C9B—C10B   | 128.6 (3) |
| C8A—C9A—H9AA   | 115.5     | C8B—C9B—H9BA   | 115.7     |
| C10A—C9A—H9AA  | 115.5     | C10B—C9B—H9BA  | 115.7     |
| C15A—C10A—C11A | 117.7 (3) | C15B—C10B—C11B | 117.3 (3) |
| C15A—C10A—C9A  | 119.3 (3) | C15B—C10B—C9B  | 119.7 (3) |
| C11A—C10A—C9A  | 123.0 (3) | C11B—C10B—C9B  | 123.0 (3) |
| C12A—C11A—C10A | 120.7 (3) | C12B—C11B—C10B | 121.3 (3) |
| C12A—C11A—H11A | 119.6     | C12B—C11B—H11B | 119.4     |
| C10A—C11A—H11A | 119.6     | C10B—C11B—H11B | 119.4     |

|                |           |                |           |
|----------------|-----------|----------------|-----------|
| C11A—C12A—C13A | 121.0 (3) | C11B—C12B—C13B | 120.5 (3) |
| C11A—C12A—H12A | 119.5     | C11B—C12B—H12B | 119.8     |
| C13A—C12A—H12A | 119.5     | C13B—C12B—H12B | 119.8     |
| O3A—C13A—C14A  | 124.9 (3) | O3B—C13B—C14B  | 125.2 (3) |
| O3A—C13A—C12A  | 115.5 (3) | O3B—C13B—C12B  | 115.0 (3) |
| C14A—C13A—C12A | 119.6 (3) | C14B—C13B—C12B | 119.8 (3) |
| C15A—C14A—C13A | 119.4 (3) | C13B—C14B—C15B | 119.0 (3) |
| C15A—C14A—H14A | 120.3     | C13B—C14B—H14B | 120.5     |
| C13A—C14A—H14A | 120.3     | C15B—C14B—H14B | 120.5     |
| C14A—C15A—C10A | 121.7 (3) | C14B—C15B—C10B | 122.2 (3) |
| C14A—C15A—H15A | 119.1     | C14B—C15B—H15B | 118.9     |
| C10A—C15A—H15A | 119.1     | C10B—C15B—H15B | 118.9     |
| O3A—C16A—C17A  | 106.4 (3) | O3B—C16B—C17B  | 106.2 (3) |
| O3A—C16A—H16A  | 110.4     | O3B—C16B—H16C  | 110.5     |
| C17A—C16A—H16A | 110.4     | C17B—C16B—H16C | 110.5     |
| O3A—C16A—H16B  | 110.4     | O3B—C16B—H16D  | 110.5     |
| C17A—C16A—H16B | 110.4     | C17B—C16B—H16D | 110.5     |
| H16A—C16A—H16B | 108.6     | H16C—C16B—H16D | 108.7     |
| C16A—C17A—C18A | 115.6 (3) | C16B—C17B—C18B | 114.6 (3) |
| C16A—C17A—H17A | 108.4     | C16B—C17B—H17C | 108.6     |
| C18A—C17A—H17A | 108.4     | C18B—C17B—H17C | 108.6     |
| C16A—C17A—H17B | 108.4     | C16B—C17B—H17D | 108.6     |
| C18A—C17A—H17B | 108.4     | C18B—C17B—H17D | 108.6     |
| H17A—C17A—H17B | 107.4     | H17C—C17B—H17D | 107.6     |
| C17A—C18A—C19A | 111.6 (3) | C17B—C18B—C19B | 111.2 (3) |
| C17A—C18A—H18A | 109.3     | C17B—C18B—H18C | 109.4     |
| C19A—C18A—H18A | 109.3     | C19B—C18B—H18C | 109.4     |
| C17A—C18A—H18B | 109.3     | C17B—C18B—H18D | 109.4     |
| C19A—C18A—H18B | 109.3     | C19B—C18B—H18D | 109.4     |
| H18A—C18A—H18B | 108.0     | H18C—C18B—H18D | 108.0     |
| C20A—C19A—C18A | 116.1 (3) | C20B—C19B—C18B | 116.2 (3) |
| C20A—C19A—H19A | 108.3     | C20B—C19B—H19C | 108.2     |
| C18A—C19A—H19A | 108.3     | C18B—C19B—H19C | 108.2     |
| C20A—C19A—H19B | 108.3     | C20B—C19B—H19D | 108.2     |
| C18A—C19A—H19B | 108.3     | C18B—C19B—H19D | 108.2     |
| H19A—C19A—H19B | 107.4     | H19C—C19B—H19D | 107.4     |
| C19A—C20A—C21A | 112.7 (3) | C19B—C20B—C21B | 112.7 (3) |
| C19A—C20A—H20A | 109.0     | C19B—C20B—H20C | 109.1     |
| C21A—C20A—H20A | 109.0     | C21B—C20B—H20C | 109.1     |
| C19A—C20A—H20B | 109.0     | C19B—C20B—H20D | 109.1     |
| C21A—C20A—H20B | 109.0     | C21B—C20B—H20D | 109.1     |
| H20A—C20A—H20B | 107.8     | H20C—C20B—H20D | 107.8     |
| C22A—C21A—C20A | 114.4 (3) | C20B—C21B—C22B | 115.0 (3) |
| C22A—C21A—H21A | 108.6     | C20B—C21B—H21C | 108.5     |
| C20A—C21A—H21A | 108.6     | C22B—C21B—H21C | 108.5     |
| C22A—C21A—H21B | 108.6     | C20B—C21B—H21D | 108.5     |
| C20A—C21A—H21B | 108.6     | C22B—C21B—H21D | 108.5     |
| H21A—C21A—H21B | 107.6     | H21C—C21B—H21D | 107.5     |

|                 |            |                 |            |
|-----------------|------------|-----------------|------------|
| C21A—C22A—C23A  | 113.3 (3)  | C23B—C22B—C21B  | 113.7 (3)  |
| C21A—C22A—H22A  | 108.9      | C23B—C22B—H22C  | 108.8      |
| C23A—C22A—H22A  | 108.9      | C21B—C22B—H22C  | 108.8      |
| C21A—C22A—H22B  | 108.9      | C23B—C22B—H22D  | 108.8      |
| C23A—C22A—H22B  | 108.9      | C21B—C22B—H22D  | 108.8      |
| H22A—C22A—H22B  | 107.7      | H22C—C22B—H22D  | 107.7      |
| C24A—C23A—C22A  | 113.7 (3)  | C22B—C23B—C24B  | 114.0 (3)  |
| C24A—C23A—H23A  | 108.8      | C22B—C23B—H23C  | 108.7      |
| C22A—C23A—H23A  | 108.8      | C24B—C23B—H23C  | 108.7      |
| C24A—C23A—H23B  | 108.8      | C22B—C23B—H23D  | 108.7      |
| C22A—C23A—H23B  | 108.8      | C24B—C23B—H23D  | 108.7      |
| H23A—C23A—H23B  | 107.7      | H23C—C23B—H23D  | 107.6      |
| C23A—C24A—C25A  | 113.9 (3)  | C23B—C24B—C25B  | 114.2 (3)  |
| C23A—C24A—H24A  | 108.8      | C23B—C24B—H24C  | 108.7      |
| C25A—C24A—H24A  | 108.8      | C25B—C24B—H24C  | 108.7      |
| C23A—C24A—H24B  | 108.8      | C23B—C24B—H24D  | 108.7      |
| C25A—C24A—H24B  | 108.8      | C25B—C24B—H24D  | 108.7      |
| H24A—C24A—H24B  | 107.7      | H24C—C24B—H24D  | 107.6      |
| C26A—C25A—C24A  | 113.7 (3)  | C26B—C25B—C24B  | 113.7 (3)  |
| C26A—C25A—H25A  | 108.8      | C26B—C25B—H25C  | 108.8      |
| C24A—C25A—H25A  | 108.8      | C24B—C25B—H25C  | 108.8      |
| C26A—C25A—H25B  | 108.8      | C26B—C25B—H25D  | 108.8      |
| C24A—C25A—H25B  | 108.8      | C24B—C25B—H25D  | 108.8      |
| H25A—C25A—H25B  | 107.7      | H25C—C25B—H25D  | 107.7      |
| C25A—C26A—C27A  | 113.5 (3)  | C25B—C26B—C27B  | 113.4 (3)  |
| C25A—C26A—H26A  | 108.9      | C25B—C26B—H26C  | 108.9      |
| C27A—C26A—H26A  | 108.9      | C27B—C26B—H26C  | 108.9      |
| C25A—C26A—H26B  | 108.9      | C25B—C26B—H26D  | 108.9      |
| C27A—C26A—H26B  | 108.9      | C27B—C26B—H26D  | 108.9      |
| H26A—C26A—H26B  | 107.7      | H26C—C26B—H26D  | 107.7      |
| C26A—C27A—H27A  | 109.5      | C26B—C27B—H27D  | 109.5      |
| C26A—C27A—H27B  | 109.5      | C26B—C27B—H27E  | 109.5      |
| H27A—C27A—H27B  | 109.5      | H27D—C27B—H27E  | 109.5      |
| C26A—C27A—H27C  | 109.5      | C26B—C27B—H27F  | 109.5      |
| H27A—C27A—H27C  | 109.5      | H27D—C27B—H27F  | 109.5      |
| H27B—C27A—H27C  | 109.5      | H27E—C27B—H27F  | 109.5      |
| <br>            |            |                 |            |
| O1A—C1A—C2A—C3A | -179.1 (3) | O1B—C1B—C2B—C3B | -179.6 (3) |
| C6A—C1A—C2A—C3A | 0.6 (5)    | C6B—C1B—C2B—C3B | -0.7 (5)   |
| C1A—C2A—C3A—C4A | 0.8 (6)    | C1B—C2B—C3B—C4B | 0.7 (6)    |
| C2A—C3A—C4A—C5A | -1.2 (6)   | C2B—C3B—C4B—C5B | -0.1 (6)   |
| C3A—C4A—C5A—C6A | 0.2 (6)    | C3B—C4B—C5B—C6B | -0.6 (6)   |
| C4A—C5A—C6A—C1A | 1.2 (5)    | C4B—C5B—C6B—C1B | 0.6 (5)    |
| C4A—C5A—C6A—C7A | -177.9 (3) | C4B—C5B—C6B—C7B | -179.4 (3) |
| O1A—C1A—C6A—C5A | 178.1 (3)  | O1B—C1B—C6B—C5B | 178.9 (3)  |
| C2A—C1A—C6A—C5A | -1.5 (5)   | C2B—C1B—C6B—C5B | 0.1 (5)    |
| O1A—C1A—C6A—C7A | -2.8 (5)   | O1B—C1B—C6B—C7B | -1.1 (5)   |
| C2A—C1A—C6A—C7A | 177.6 (3)  | C2B—C1B—C6B—C7B | 180.0 (3)  |

|                     |            |                     |            |
|---------------------|------------|---------------------|------------|
| C5A—C6A—C7A—O2A     | 178.4 (3)  | C5B—C6B—C7B—O2B     | 175.9 (3)  |
| C1A—C6A—C7A—O2A     | -0.7 (5)   | C1B—C6B—C7B—O2B     | -4.1 (5)   |
| C5A—C6A—C7A—C8A     | -2.7 (5)   | C5B—C6B—C7B—C8B     | -5.4 (5)   |
| C1A—C6A—C7A—C8A     | 178.2 (3)  | C1B—C6B—C7B—C8B     | 174.7 (3)  |
| O2A—C7A—C8A—C9A     | 6.5 (5)    | O2B—C7B—C8B—C9B     | 8.8 (5)    |
| C6A—C7A—C8A—C9A     | -172.3 (3) | C6B—C7B—C8B—C9B     | -169.9 (3) |
| C7A—C8A—C9A—C10A    | 179.9 (3)  | C7B—C8B—C9B—C10B    | -179.7 (3) |
| C8A—C9A—C10A—C15A   | -168.6 (4) | C8B—C9B—C10B—C15B   | -169.5 (4) |
| C8A—C9A—C10A—C11A   | 13.0 (6)   | C8B—C9B—C10B—C11B   | 10.8 (6)   |
| C15A—C10A—C11A—C12A | 0.1 (5)    | C15B—C10B—C11B—C12B | -0.2 (5)   |
| C9A—C10A—C11A—C12A  | 178.5 (3)  | C9B—C10B—C11B—C12B  | 179.5 (3)  |
| C10A—C11A—C12A—C13A | 0.2 (5)    | C10B—C11B—C12B—C13B | 0.3 (5)    |
| C16A—O3A—C13A—C14A  | -6.8 (5)   | C16B—O3B—C13B—C14B  | -12.2 (5)  |
| C16A—O3A—C13A—C12A  | 173.7 (3)  | C16B—O3B—C13B—C12B  | 168.4 (3)  |
| C11A—C12A—C13A—O3A  | 179.8 (3)  | C11B—C12B—C13B—O3B  | 178.9 (3)  |
| C11A—C12A—C13A—C14A | 0.3 (5)    | C11B—C12B—C13B—C14B | -0.5 (5)   |
| O3A—C13A—C14A—C15A  | 179.5 (3)  | O3B—C13B—C14B—C15B  | -178.6 (3) |
| C12A—C13A—C14A—C15A | -1.0 (5)   | C12B—C13B—C14B—C15B | 0.8 (5)    |
| C13A—C14A—C15A—C10A | 1.4 (5)    | C13B—C14B—C15B—C10B | -0.7 (5)   |
| C11A—C10A—C15A—C14A | -0.9 (5)   | C11B—C10B—C15B—C14B | 0.5 (5)    |
| C9A—C10A—C15A—C14A  | -179.4 (3) | C9B—C10B—C15B—C14B  | -179.2 (3) |
| C13A—O3A—C16A—C17A  | 175.0 (3)  | C13B—O3B—C16B—C17B  | 177.5 (3)  |
| O3A—C16A—C17A—C18A  | 176.8 (3)  | O3B—C16B—C17B—C18B  | 176.5 (3)  |
| C16A—C17A—C18A—C19A | -176.9 (3) | C16B—C17B—C18B—C19B | -178.8 (3) |
| C17A—C18A—C19A—C20A | -176.2 (3) | C17B—C18B—C19B—C20B | -173.5 (3) |
| C18A—C19A—C20A—C21A | -179.7 (3) | C18B—C19B—C20B—C21B | -176.4 (3) |
| C19A—C20A—C21A—C22A | -174.4 (3) | C19B—C20B—C21B—C22B | -174.4 (3) |
| C20A—C21A—C22A—C23A | 175.3 (3)  | C20B—C21B—C22B—C23B | 177.0 (3)  |
| C21A—C22A—C23A—C24A | -177.5 (3) | C21B—C22B—C23B—C24B | -178.3 (3) |
| C22A—C23A—C24A—C25A | 176.3 (3)  | C22B—C23B—C24B—C25B | 173.9 (3)  |
| C23A—C24A—C25A—C26A | 179.6 (3)  | C23B—C24B—C25B—C26B | 178.6 (3)  |
| C24A—C25A—C26A—C27A | -179.5 (3) | C24B—C25B—C26B—C27B | -174.1 (3) |

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

| $D\text{—H}\cdots A$ | $D\text{—H}$ | $H\cdots A$ | $D\cdots A$ | $D\text{—H}\cdots A$ |
|----------------------|--------------|-------------|-------------|----------------------|
| O1A—H1A…O2A          | 0.82         | 1.79        | 2.513 (4)   | 146                  |
| O1B—H1B…O2B          | 0.82         | 1.81        | 2.530 (4)   | 146                  |
| C22B—H22C… $Cg1^i$   | 0.97         | 2.77        | 3.654 (4)   | 151                  |
| C16B—H16D… $Cg2^i$   | 0.97         | 3.00        | 3.736 (4)   | 134                  |
| C17B—H17D… $Cg2$     | 0.97         | 2.82        | 3.612 (4)   | 139                  |
| C22A—H22B… $Cg3$     | 0.97         | 2.93        | 3.765 (4)   | 145                  |

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