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10,10-Dimethylanthrone

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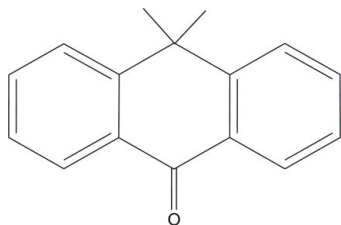
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Key indicators: single-crystal X-ray study; $T = 100$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; R factor = 0.044; wR factor = 0.142; data-to-parameter ratio = 21.5.

In the title compound, $\text{C}_{16}\text{H}_{14}\text{O}$, the asymmetric unit consists of three crystallographically independent molecules. The anthracene units are essentially planar, with maximum deviations of 0.165 (1), 0.153 (1) and 0.045 (1) Å in the three molecules. In the crystal structure, molecules are linked *via* intermolecular $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds. Further stabilization is provided by $\text{C}-\text{H}\cdots\pi$ interactions.

Related literature

For analytical applications of the title compound, see: Trevelyan (1952). For related structures, see: Destro *et al.* (1973); Ghosh *et al.* (1993); Iball & Low (1974); Srivastava (1964); Zhou *et al.* (2004; 2005; 2007). For bond-length data, see: Allen *et al.* (1987). For the stability of the temperature controller used in the data collection, see: Cosier & Glazer (1986).



Experimental

Crystal data

$\text{C}_{16}\text{H}_{14}\text{O}$	$\alpha = 107.955$ (1)°
$M_r = 222.27$	$\beta = 98.734$ (1)°
Triclinic, $P\bar{1}$	$\gamma = 111.764$ (1)°
$a = 11.2438$ (6) Å	$V = 1732.47$ (16) Å ³
$b = 12.1105$ (6) Å	$Z = 6$
$c = 15.1025$ (8) Å	Mo $K\alpha$ radiation

† Thomson Reuters ResearcherID: A-3561-2009.

 $\mu = 0.08$ mm⁻¹
 $T = 100$ K

0.62 × 0.31 × 0.27 mm

Data collection

 Bruker APEX DUO CCD area-detector diffractometer
 Absorption correction: multi-scan (SADABS; Bruker, 2009)
 $T_{\min} = 0.953$, $T_{\max} = 0.979$

 35076 measured reflections
 10016 independent reflections
 8420 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.028$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.044$
 $wR(F^2) = 0.142$
 $S = 1.07$
 10016 reflections

 466 parameters
 H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.54$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.36$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$\text{Cg}1$, $\text{Cg}7$, $\text{Cg}13$ and $\text{Cg}15$ are the centroids of the $\text{C}1\text{A}-\text{C}6\text{A}$, $\text{C}1\text{B}-\text{C}6\text{B}$, $\text{C}1\text{C}-\text{C}6\text{C}$ and $\text{C}8\text{C}-\text{C}13\text{C}$ rings, respectively.

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{C}10\text{A}-\text{H}10\text{A}\cdots\text{O}1\text{C}^{\text{i}}$	0.93	2.52	3.4097 (16)	160
$\text{C}10\text{C}-\text{H}10\text{C}\cdots\text{O}1\text{B}^{\text{ii}}$	0.93	2.59	3.4374 (18)	151
$\text{C}3\text{C}-\text{H}3\text{CA}\cdots\text{O}1\text{A}^{\text{iii}}$	0.93	2.45	3.2287 (17)	141
$\text{C}15\text{B}-\text{H}15\text{D}\cdots\text{O}1\text{B}^{\text{ii}}$	0.96	2.43	3.3568 (18)	161
$\text{C}16\text{A}-\text{H}16\text{A}\cdots\text{O}1\text{A}^{\text{iv}}$	0.96	2.48	3.3934 (16)	159
$\text{C}3\text{A}-\text{H}3\text{AA}\cdots\text{Cg}13^{\text{v}}$	0.93	2.69	3.3996 (14)	134
$\text{C}3\text{B}-\text{H}3\text{BA}\cdots\text{Cg}13^{\text{vi}}$	0.93	2.77	3.3715 (14)	123
$\text{C}15\text{A}-\text{H}15\text{B}\cdots\text{Cg}15^{\text{v}}$	0.96	2.71	3.4915 (15)	139
$\text{C}15\text{C}-\text{H}15\text{G}\cdots\text{Cg}1^{\text{vii}}$	0.96	2.96	3.8656 (13)	158
$\text{C}16\text{C}-\text{H}16\text{G}\cdots\text{Cg}7^{\text{vi}}$	0.96	2.91	3.7432 (14)	145

Symmetry codes: (i) $x - 1, y, z$; (ii) $-x + 1, -y + 1, -z$; (iii) $-x + 2, -y + 1, -z + 1$; (iv) $-x + 1, -y + 2, -z + 1$; (v) $x, y + 1, z$; (vi) $-x + 2, -y + 1, -z$; (vii) $x, y - 1, z$.

Data collection: APEX2 (Bruker, 2009); cell refinement: SAINT (Bruker, 2009); data reduction: SAINT; program(s) used to solve structure: SHELXTL (Sheldrick, 2008); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL and PLATON (Spek, 2009).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: SJ2742).

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supporting information

Acta Cryst. (2010). E66, o808–o809 [doi:10.1107/S1600536810008524]

10,10-Dimethylantrone

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S1. Comment

Anthrone is a tricyclic aromatic hydrocarbon which is used for a popular cellulose assay and in the colorimetric determination of carbohydrates (Trevelyan, 1952). The crystal structures of anthrone (Srivastava, 1964), 10-bromoanthrone (Destro *et al.*, 1973), 9,10-dimethylantracene (Iball & Low, 1974), benzylideneanthrone at 193 K (Ghosh *et al.*, 1993), 10-(2-methylbenzylidene)anthrone (Zhou *et al.*, 2004), 10-(3,4-dimethoxybenzylidene) anthrone (Zhou *et al.*, 2005) and 10-(4-hydroxy-3-nitrobenzylidene) anthrone (Zhou *et al.*, 2007), have been reported. In view of the importance of the title compound, (I), its crystal structure is reported here.

The asymmetric unit of the title compound, consists of three crystallographically independent 10,10-Dimethylantrone molecules, (A, B & C), as shown in Fig. 1. The bond lengths and angles of molecules A, B and C agree with each other and are within normal ranges (Allen *et al.*, 1987). The anthracene units are essentially planar with maximum deviations of 0.165 (1)Å for atom C3A (molecule A), 0.153 (1)Å for atom C14B (molecule B) and 0.045 (1)Å for atom C14C (molecule C). The two fused benzene rings, C1–C6 and C8–C13, make dihedral angles with the C1/C6–C8/C13–C14 plane of 5.74 (5)° and 3.85 (5)° in molecule A; 4.40 (6)° and 2.79 (6)° in molecule B; and 1.69 (6)° and 1.63 (6)° in molecule C.

In the crystal structure (Fig. 2), the molecules are linked through intermolecular C10—H10A···O1C, C10C—H10C···O1B, C3C—H3CA···O1A C15B—H15D···O1B and C16A—H16A···O1A hydrogen bonds. The structure is further stabilized by C—H··· π interactions (Table 1), involving C1A–C6A (centroid Cg1), C1B–C6B (centroid Cg7), C1C–C6C (centroid Cg13) and C8C–C13C (centroid Cg15).

S2. Experimental

The title compound was obtained as a gift sample from R. L. Fine Chem, Bangalore, India. X-ray quality crystals were obtained by slow evaporation from a methanol solution (m. p.: 349–352 K).

S3. Refinement

All hydrogen atoms were positioned geometrically [C–H = 0.93 or 0.96Å] and were refined using a riding model, with $U_{\text{iso}}(\text{H}) = 1.2$ or $1.5 U_{\text{eq}}(\text{C})$. A rotating group model was applied to the methyl groups.

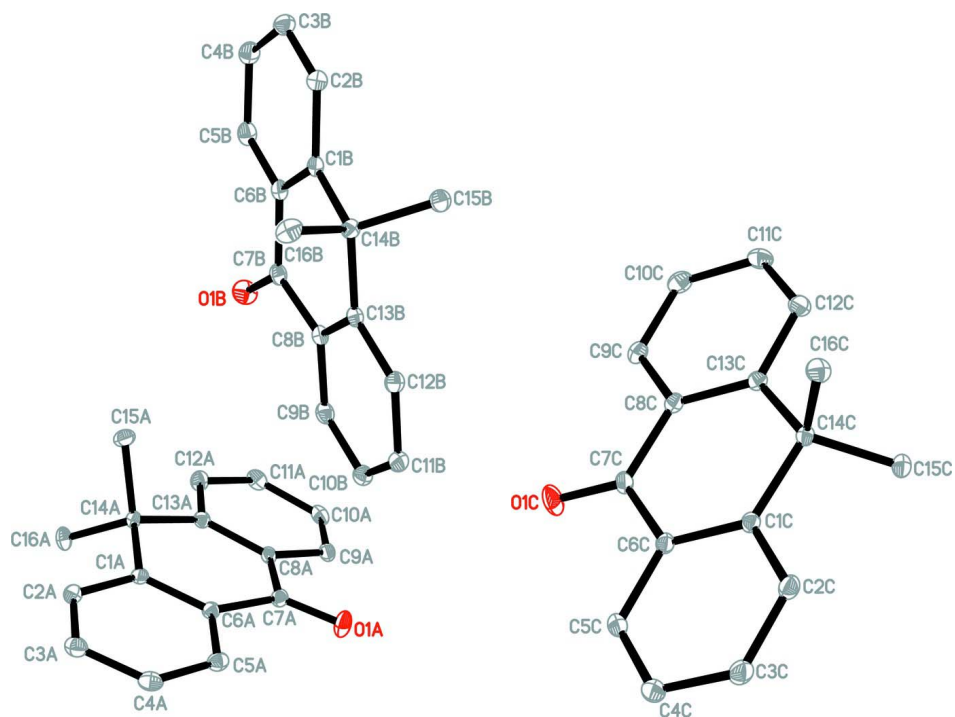


Figure 1

The asymmetric unit of the title compound, showing 30% probability displacement ellipsoids and the atom-numbering scheme. H atoms are omitted for clarity.

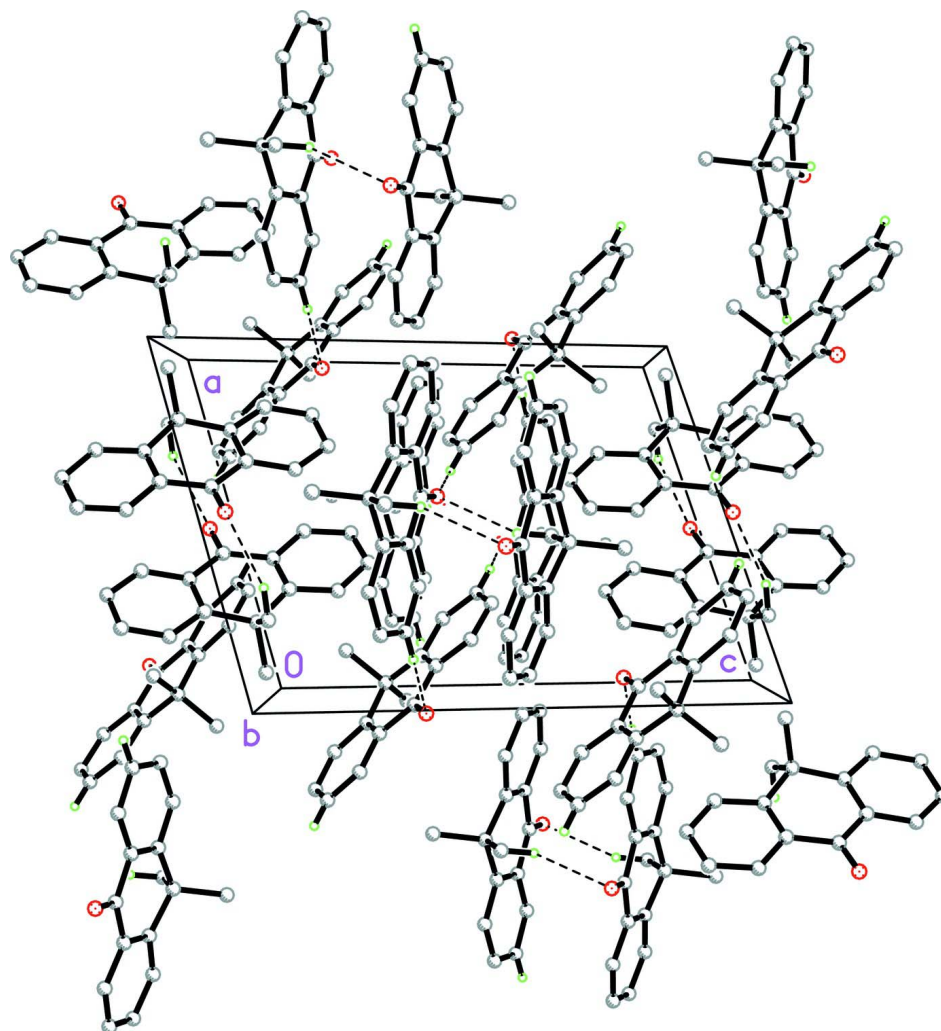


Figure 2

Crystal packing of the title compound, showing the hydrogen-bonding network (dashed lines). H atoms not involved in the hydrogen bond interactions are omitted for clarity.

10,10-Dimethyl-10*H*-anthracen-9-one

Crystal data

$C_{16}H_{14}O$

$M_r = 222.27$

Triclinic, $P\bar{1}$

Hall symbol: $-P\ 1$

$a = 11.2438\ (6)\ \text{\AA}$

$b = 12.1105\ (6)\ \text{\AA}$

$c = 15.1025\ (8)\ \text{\AA}$

$\alpha = 107.955\ (1)^\circ$

$\beta = 98.734\ (1)^\circ$

$\gamma = 111.764\ (1)^\circ$

$V = 1732.47\ (16)\ \text{\AA}^3$

$Z = 6$

$F(000) = 708$

$D_x = 1.278\ \text{Mg m}^{-3}$

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

Cell parameters from 9906 reflections

$\theta = 2.8\text{--}32.7^\circ$

$\mu = 0.08\ \text{mm}^{-1}$

$T = 100\ \text{K}$

Block, orange

$0.62 \times 0.31 \times 0.27\ \text{mm}$

Data collection

Bruker APEX DUO CCD area-detector
diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
 φ and ω scans
Absorption correction: multi-scan
(*SADABS*; Bruker, 2009)
 $T_{\min} = 0.953$, $T_{\max} = 0.979$

35076 measured reflections
10016 independent reflections
8420 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.028$
 $\theta_{\text{max}} = 30.0^\circ$, $\theta_{\text{min}} = 1.5^\circ$
 $h = -15 \rightarrow 15$
 $k = -17 \rightarrow 17$
 $l = -21 \rightarrow 21$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.044$
 $wR(F^2) = 0.142$
 $S = 1.07$
10016 reflections
466 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
 $w = 1/[\sigma^2(F_o^2) + (0.0807P)^2 + 0.476P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.001$
 $\Delta\rho_{\text{max}} = 0.54 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.36 \text{ e } \text{\AA}^{-3}$

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 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>	$U_{\text{iso}}^*/U_{\text{eq}}$
O1A	0.56816 (9)	0.79013 (8)	0.44954 (7)	0.02699 (19)
C1A	0.68614 (10)	1.07682 (10)	0.39587 (7)	0.01479 (19)
C2A	0.80724 (11)	1.18655 (11)	0.41974 (8)	0.0192 (2)
H2AA	0.8086	1.2474	0.3940	0.023*
C3A	0.92550 (11)	1.20656 (11)	0.48111 (8)	0.0218 (2)
H3AA	1.0040	1.2814	0.4972	0.026*
C4A	0.92729 (11)	1.11539 (12)	0.51856 (8)	0.0218 (2)
H4AA	1.0068	1.1278	0.5584	0.026*
C5A	0.80939 (11)	1.00627 (11)	0.49582 (8)	0.0194 (2)
H5AA	0.8097	0.9447	0.5203	0.023*
C6A	0.68883 (10)	0.98739 (10)	0.43599 (7)	0.01541 (19)
C7A	0.56463 (11)	0.87236 (10)	0.41915 (7)	0.01671 (19)
C8A	0.43582 (10)	0.86035 (10)	0.36546 (7)	0.01535 (19)
C9A	0.31622 (11)	0.75993 (10)	0.35921 (8)	0.0189 (2)

H9AA	0.3201	0.7045	0.3894	0.023*
C10A	0.19306 (11)	0.74253 (10)	0.30882 (8)	0.0210 (2)
H10A	0.1141	0.6768	0.3058	0.025*
C11A	0.18879 (11)	0.82504 (11)	0.26257 (9)	0.0223 (2)
H11A	0.1065	0.8137	0.2278	0.027*
C12A	0.30651 (11)	0.92392 (11)	0.26805 (8)	0.0209 (2)
H12A	0.3018	0.9771	0.2358	0.025*
C13A	0.43266 (10)	0.94566 (10)	0.32106 (7)	0.01543 (19)
C14A	0.55932 (10)	1.05499 (10)	0.32446 (7)	0.01520 (19)
C15A	0.57639 (12)	1.01805 (12)	0.22108 (8)	0.0218 (2)
H15A	0.5831	0.9378	0.2022	0.033*
H15B	0.6565	1.0850	0.2215	0.033*
H15C	0.5001	1.0085	0.1754	0.033*
C16A	0.54310 (12)	1.18059 (10)	0.35169 (9)	0.0216 (2)
H16A	0.5360	1.2064	0.4167	0.032*
H16B	0.4634	1.1662	0.3068	0.032*
H16C	0.6198	1.2477	0.3486	0.032*
O1B	0.52412 (9)	0.74204 (9)	0.03541 (7)	0.02682 (19)
C1B	0.69616 (10)	0.58076 (10)	-0.09660 (7)	0.01525 (19)
C2B	0.70059 (11)	0.53353 (11)	-0.19275 (8)	0.0196 (2)
H2BA	0.7581	0.4961	-0.2062	0.023*
C3B	0.62091 (12)	0.54135 (11)	-0.26839 (8)	0.0226 (2)
H3BA	0.6242	0.5077	-0.3318	0.027*
C4B	0.53651 (12)	0.59936 (11)	-0.24945 (8)	0.0238 (2)
H4BA	0.4842	0.6057	-0.2998	0.029*
C5B	0.53076 (11)	0.64754 (11)	-0.15537 (8)	0.0210 (2)
H5BA	0.4746	0.6868	-0.1424	0.025*
C6B	0.60928 (10)	0.63766 (10)	-0.07907 (8)	0.01640 (19)
C7B	0.59492 (10)	0.68598 (10)	0.01946 (8)	0.01716 (19)
C8B	0.66776 (10)	0.66388 (10)	0.09737 (8)	0.01560 (19)
C9B	0.64508 (11)	0.69874 (10)	0.18895 (8)	0.0186 (2)
H9BA	0.5841	0.7329	0.1983	0.022*
C10B	0.71228 (12)	0.68289 (11)	0.26493 (8)	0.0203 (2)
H10B	0.6968	0.7061	0.3252	0.024*
C11B	0.80371 (11)	0.63168 (11)	0.25022 (8)	0.0195 (2)
H11B	0.8496	0.6208	0.3010	0.023*
C12B	0.82647 (11)	0.59700 (10)	0.16019 (8)	0.0178 (2)
H12B	0.8880	0.5634	0.1517	0.021*
C13B	0.75874 (10)	0.61140 (9)	0.08158 (7)	0.01495 (18)
C14B	0.78902 (10)	0.57404 (10)	-0.01569 (7)	0.01553 (19)
C15B	0.77900 (13)	0.43618 (11)	-0.04709 (8)	0.0236 (2)
H15D	0.6880	0.3751	-0.0599	0.035*
H15E	0.8370	0.4322	0.0042	0.035*
H15F	0.8057	0.4155	-0.1051	0.035*
C16B	0.93459 (11)	0.67118 (12)	-0.00016 (9)	0.0234 (2)
H16D	0.9418	0.7573	0.0205	0.035*
H16E	0.9563	0.6495	-0.0603	0.035*
H16F	0.9958	0.6669	0.0490	0.035*

O1C	0.95232 (9)	0.45834 (8)	0.29323 (7)	0.0297 (2)
C1C	1.10277 (10)	0.24592 (9)	0.31380 (7)	0.01459 (18)
C2C	1.20914 (11)	0.25001 (10)	0.37943 (8)	0.0185 (2)
H2CA	1.2281	0.1791	0.3644	0.022*
C3C	1.28635 (12)	0.35724 (11)	0.46597 (8)	0.0230 (2)
H3CA	1.3558	0.3574	0.5083	0.028*
C4C	1.26000 (13)	0.46478 (11)	0.48958 (8)	0.0247 (2)
H4CA	1.3111	0.5365	0.5479	0.030*
C5C	1.15738 (12)	0.46413 (10)	0.42563 (8)	0.0201 (2)
H5CA	1.1402	0.5361	0.4408	0.024*
C6C	1.07904 (10)	0.35551 (9)	0.33796 (7)	0.01501 (19)
C7C	0.97047 (10)	0.36071 (10)	0.27270 (8)	0.01652 (19)
C8C	0.88406 (10)	0.24447 (10)	0.18236 (7)	0.01471 (18)
C9C	0.77830 (11)	0.24772 (11)	0.12168 (8)	0.0180 (2)
H9CA	0.7660	0.3225	0.1388	0.022*
C10C	0.69261 (11)	0.14160 (12)	0.03723 (8)	0.0219 (2)
H10C	0.6232	0.1445	-0.0028	0.026*
C11C	0.71192 (12)	0.03012 (11)	0.01295 (8)	0.0226 (2)
H11C	0.6544	-0.0424	-0.0434	0.027*
C12C	0.81653 (11)	0.02660 (11)	0.07228 (8)	0.0193 (2)
H12C	0.8282	-0.0485	0.0545	0.023*
C13C	0.90527 (10)	0.13340 (10)	0.15846 (7)	0.01460 (18)
C14C	1.01912 (10)	0.12450 (10)	0.22050 (7)	0.01462 (18)
C15C	0.95875 (11)	0.00734 (10)	0.24858 (8)	0.0194 (2)
H15G	0.8980	0.0188	0.2848	0.029*
H15H	1.0297	0.0013	0.2880	0.029*
H15I	0.9111	-0.0707	0.1904	0.029*
C16C	1.11272 (11)	0.10352 (11)	0.15848 (8)	0.0197 (2)
H16G	1.1483	0.1751	0.1397	0.030*
H16H	1.0627	0.0248	0.1011	0.030*
H16I	1.1853	0.0977	0.1962	0.030*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1A	0.0275 (4)	0.0230 (4)	0.0374 (5)	0.0127 (4)	0.0079 (4)	0.0198 (4)
C1A	0.0167 (5)	0.0150 (4)	0.0130 (4)	0.0075 (4)	0.0056 (3)	0.0051 (3)
C2A	0.0200 (5)	0.0183 (5)	0.0180 (5)	0.0061 (4)	0.0066 (4)	0.0081 (4)
C3A	0.0170 (5)	0.0229 (5)	0.0190 (5)	0.0043 (4)	0.0056 (4)	0.0058 (4)
C4A	0.0184 (5)	0.0271 (6)	0.0172 (5)	0.0105 (4)	0.0033 (4)	0.0058 (4)
C5A	0.0209 (5)	0.0228 (5)	0.0177 (5)	0.0127 (4)	0.0052 (4)	0.0086 (4)
C6A	0.0175 (5)	0.0157 (4)	0.0144 (4)	0.0087 (4)	0.0055 (3)	0.0056 (4)
C7A	0.0194 (5)	0.0157 (4)	0.0175 (4)	0.0092 (4)	0.0065 (4)	0.0074 (4)
C8A	0.0180 (5)	0.0129 (4)	0.0158 (4)	0.0078 (4)	0.0061 (3)	0.0049 (3)
C9A	0.0222 (5)	0.0138 (4)	0.0201 (5)	0.0069 (4)	0.0087 (4)	0.0062 (4)
C10A	0.0184 (5)	0.0155 (5)	0.0231 (5)	0.0048 (4)	0.0073 (4)	0.0031 (4)
C11A	0.0163 (5)	0.0212 (5)	0.0250 (5)	0.0082 (4)	0.0035 (4)	0.0050 (4)
C12A	0.0197 (5)	0.0203 (5)	0.0243 (5)	0.0103 (4)	0.0049 (4)	0.0100 (4)

C13A	0.0167 (5)	0.0143 (4)	0.0163 (4)	0.0081 (4)	0.0057 (4)	0.0054 (4)
C14A	0.0174 (5)	0.0140 (4)	0.0154 (4)	0.0071 (4)	0.0051 (3)	0.0071 (3)
C15A	0.0227 (5)	0.0260 (5)	0.0160 (5)	0.0088 (4)	0.0062 (4)	0.0098 (4)
C16A	0.0231 (5)	0.0137 (5)	0.0302 (6)	0.0101 (4)	0.0077 (4)	0.0091 (4)
O1B	0.0291 (4)	0.0303 (5)	0.0296 (4)	0.0230 (4)	0.0088 (3)	0.0105 (4)
C1B	0.0144 (4)	0.0130 (4)	0.0182 (4)	0.0055 (4)	0.0038 (3)	0.0073 (4)
C2B	0.0201 (5)	0.0191 (5)	0.0195 (5)	0.0086 (4)	0.0060 (4)	0.0078 (4)
C3B	0.0231 (5)	0.0219 (5)	0.0181 (5)	0.0060 (4)	0.0034 (4)	0.0084 (4)
C4B	0.0216 (5)	0.0219 (5)	0.0217 (5)	0.0059 (4)	−0.0016 (4)	0.0096 (4)
C5B	0.0174 (5)	0.0182 (5)	0.0257 (5)	0.0078 (4)	0.0014 (4)	0.0091 (4)
C6B	0.0146 (4)	0.0130 (4)	0.0205 (5)	0.0058 (4)	0.0030 (4)	0.0066 (4)
C7B	0.0159 (5)	0.0145 (4)	0.0215 (5)	0.0078 (4)	0.0046 (4)	0.0069 (4)
C8B	0.0146 (4)	0.0124 (4)	0.0193 (5)	0.0058 (4)	0.0051 (4)	0.0058 (4)
C9B	0.0189 (5)	0.0156 (5)	0.0227 (5)	0.0089 (4)	0.0087 (4)	0.0067 (4)
C10B	0.0241 (5)	0.0176 (5)	0.0193 (5)	0.0088 (4)	0.0092 (4)	0.0068 (4)
C11B	0.0204 (5)	0.0189 (5)	0.0184 (5)	0.0079 (4)	0.0044 (4)	0.0080 (4)
C12B	0.0169 (5)	0.0177 (5)	0.0207 (5)	0.0090 (4)	0.0057 (4)	0.0084 (4)
C13B	0.0137 (4)	0.0124 (4)	0.0178 (4)	0.0050 (3)	0.0050 (3)	0.0056 (3)
C14B	0.0157 (4)	0.0165 (4)	0.0176 (4)	0.0095 (4)	0.0059 (3)	0.0076 (4)
C15B	0.0354 (6)	0.0232 (5)	0.0222 (5)	0.0209 (5)	0.0113 (5)	0.0106 (4)
C16B	0.0154 (5)	0.0323 (6)	0.0236 (5)	0.0090 (4)	0.0069 (4)	0.0139 (5)
O1C	0.0302 (5)	0.0166 (4)	0.0363 (5)	0.0140 (4)	−0.0027 (4)	0.0047 (3)
C1C	0.0147 (4)	0.0139 (4)	0.0151 (4)	0.0053 (4)	0.0041 (3)	0.0071 (3)
C2C	0.0186 (5)	0.0176 (5)	0.0204 (5)	0.0084 (4)	0.0032 (4)	0.0097 (4)
C3C	0.0217 (5)	0.0213 (5)	0.0212 (5)	0.0069 (4)	−0.0018 (4)	0.0096 (4)
C4C	0.0271 (6)	0.0174 (5)	0.0193 (5)	0.0053 (4)	−0.0030 (4)	0.0048 (4)
C5C	0.0231 (5)	0.0137 (4)	0.0196 (5)	0.0068 (4)	0.0024 (4)	0.0056 (4)
C6C	0.0155 (4)	0.0128 (4)	0.0159 (4)	0.0052 (4)	0.0034 (3)	0.0066 (4)
C7C	0.0162 (5)	0.0130 (4)	0.0193 (5)	0.0058 (4)	0.0033 (4)	0.0069 (4)
C8C	0.0141 (4)	0.0150 (4)	0.0156 (4)	0.0058 (4)	0.0043 (3)	0.0077 (4)
C9C	0.0172 (5)	0.0199 (5)	0.0205 (5)	0.0092 (4)	0.0051 (4)	0.0114 (4)
C10C	0.0185 (5)	0.0279 (6)	0.0196 (5)	0.0102 (4)	0.0023 (4)	0.0114 (4)
C11C	0.0210 (5)	0.0235 (5)	0.0156 (5)	0.0070 (4)	0.0005 (4)	0.0039 (4)
C12C	0.0205 (5)	0.0183 (5)	0.0170 (5)	0.0089 (4)	0.0043 (4)	0.0047 (4)
C13C	0.0146 (4)	0.0155 (4)	0.0143 (4)	0.0063 (4)	0.0049 (3)	0.0069 (4)
C14C	0.0162 (4)	0.0134 (4)	0.0151 (4)	0.0075 (4)	0.0040 (3)	0.0061 (3)
C15C	0.0229 (5)	0.0142 (4)	0.0201 (5)	0.0072 (4)	0.0046 (4)	0.0080 (4)
C16C	0.0195 (5)	0.0222 (5)	0.0205 (5)	0.0123 (4)	0.0072 (4)	0.0079 (4)

Geometric parameters (Å, °)

O1A—C7A	1.2286 (13)	C9B—H9BA	0.9300
C1A—C2A	1.4013 (14)	C10B—C11B	1.3957 (16)
C1A—C6A	1.4016 (14)	C10B—H10B	0.9300
C1A—C14A	1.5309 (14)	C11B—C12B	1.3871 (15)
C2A—C3A	1.3908 (16)	C11B—H11B	0.9300
C2A—H2AA	0.9300	C12B—C13B	1.4040 (14)
C3A—C4A	1.3920 (17)	C12B—H12B	0.9300

C3A—H3AA	0.9300	C13B—C14B	1.5281 (14)
C4A—C5A	1.3800 (16)	C14B—C15B	1.5430 (15)
C4A—H4AA	0.9300	C14B—C16B	1.5503 (15)
C5A—C6A	1.4061 (14)	C15B—H15D	0.9600
C5A—H5AA	0.9300	C15B—H15E	0.9600
C6A—C7A	1.4787 (14)	C15B—H15F	0.9600
C7A—C8A	1.4800 (15)	C16B—H16D	0.9600
C8A—C13A	1.4020 (14)	C16B—H16E	0.9600
C8A—C9A	1.4051 (14)	C16B—H16F	0.9600
C9A—C10A	1.3804 (16)	O1C—C7C	1.2280 (13)
C9A—H9AA	0.9300	C1C—C6C	1.3998 (14)
C10A—C11A	1.3945 (17)	C1C—C2C	1.4075 (14)
C10A—H10A	0.9300	C1C—C14C	1.5254 (14)
C11A—C12A	1.3867 (16)	C2C—C3C	1.3861 (15)
C11A—H11A	0.9300	C2C—H2CA	0.9300
C12A—C13A	1.4037 (15)	C3C—C4C	1.3939 (17)
C12A—H12A	0.9300	C3C—H3CA	0.9300
C13A—C14A	1.5235 (14)	C4C—C5C	1.3836 (16)
C14A—C16A	1.5359 (14)	C4C—H4CA	0.9300
C14A—C15A	1.5487 (14)	C5C—C6C	1.4043 (14)
C15A—H15A	0.9600	C5C—H5CA	0.9300
C15A—H15B	0.9600	C6C—C7C	1.4818 (14)
C15A—H15C	0.9600	C7C—C8C	1.4786 (14)
C16A—H16A	0.9600	C8C—C13C	1.4004 (14)
C16A—H16B	0.9600	C8C—C9C	1.4076 (14)
C16A—H16C	0.9600	C9C—C10C	1.3800 (15)
O1B—C7B	1.2292 (13)	C9C—H9CA	0.9300
C1B—C6B	1.4015 (14)	C10C—C11C	1.3931 (17)
C1B—C2B	1.4032 (14)	C10C—H10C	0.9300
C1B—C14B	1.5254 (14)	C11C—C12C	1.3889 (15)
C2B—C3B	1.3912 (15)	C11C—H11C	0.9300
C2B—H2BA	0.9300	C12C—C13C	1.4047 (14)
C3B—C4B	1.3889 (17)	C12C—H12C	0.9300
C3B—H3BA	0.9300	C13C—C14C	1.5260 (14)
C4B—C5B	1.3814 (17)	C14C—C15C	1.5485 (14)
C4B—H4BA	0.9300	C14C—C16C	1.5494 (14)
C5B—C6B	1.4058 (14)	C15C—H15G	0.9600
C5B—H5BA	0.9300	C15C—H15H	0.9600
C6B—C7B	1.4795 (15)	C15C—H15I	0.9600
C7B—C8B	1.4801 (14)	C16C—H16G	0.9600
C8B—C13B	1.4035 (14)	C16C—H16H	0.9600
C8B—C9B	1.4082 (14)	C16C—H16I	0.9600
C9B—C10B	1.3801 (16)		
C2A—C1A—C6A	117.35 (10)	C11B—C10B—H10B	120.4
C2A—C1A—C14A	120.03 (9)	C12B—C11B—C10B	120.30 (10)
C6A—C1A—C14A	122.55 (9)	C12B—C11B—H11B	119.9
C3A—C2A—C1A	121.50 (10)	C10B—C11B—H11B	119.9

C3A—C2A—H2AA	119.3	C11B—C12B—C13B	121.56 (10)
C1A—C2A—H2AA	119.3	C11B—C12B—H12B	119.2
C2A—C3A—C4A	120.49 (10)	C13B—C12B—H12B	119.2
C2A—C3A—H3AA	119.8	C8B—C13B—C12B	117.68 (9)
C4A—C3A—H3AA	119.8	C8B—C13B—C14B	122.66 (9)
C5A—C4A—C3A	119.08 (10)	C12B—C13B—C14B	119.63 (9)
C5A—C4A—H4AA	120.5	C1B—C14B—C13B	113.08 (8)
C3A—C4A—H4AA	120.5	C1B—C14B—C15B	109.74 (9)
C4A—C5A—C6A	120.61 (10)	C13B—C14B—C15B	109.38 (8)
C4A—C5A—H5AA	119.7	C1B—C14B—C16B	107.61 (8)
C6A—C5A—H5AA	119.7	C13B—C14B—C16B	107.82 (8)
C1A—C6A—C5A	120.93 (10)	C15B—C14B—C16B	109.10 (9)
C1A—C6A—C7A	121.15 (9)	C14B—C15B—H15D	109.5
C5A—C6A—C7A	117.90 (9)	C14B—C15B—H15E	109.5
O1A—C7A—C6A	121.08 (10)	H15D—C15B—H15E	109.5
O1A—C7A—C8A	121.12 (10)	C14B—C15B—H15F	109.5
C6A—C7A—C8A	117.80 (9)	H15D—C15B—H15F	109.5
C13A—C8A—C9A	120.68 (10)	H15E—C15B—H15F	109.5
C13A—C8A—C7A	121.17 (9)	C14B—C16B—H16D	109.5
C9A—C8A—C7A	118.15 (9)	C14B—C16B—H16E	109.5
C10A—C9A—C8A	120.83 (10)	H16D—C16B—H16E	109.5
C10A—C9A—H9AA	119.6	C14B—C16B—H16F	109.5
C8A—C9A—H9AA	119.6	H16D—C16B—H16F	109.5
C9A—C10A—C11A	119.01 (10)	H16E—C16B—H16F	109.5
C9A—C10A—H10A	120.5	C6C—C1C—C2C	117.59 (9)
C11A—C10A—H10A	120.5	C6C—C1C—C14C	122.92 (9)
C12A—C11A—C10A	120.43 (10)	C2C—C1C—C14C	119.48 (9)
C12A—C11A—H11A	119.8	C3C—C2C—C1C	121.57 (10)
C10A—C11A—H11A	119.8	C3C—C2C—H2CA	119.2
C11A—C12A—C13A	121.57 (10)	C1C—C2C—H2CA	119.2
C11A—C12A—H12A	119.2	C2C—C3C—C4C	120.06 (10)
C13A—C12A—H12A	119.2	C2C—C3C—H3CA	120.0
C8A—C13A—C12A	117.44 (10)	C4C—C3C—H3CA	120.0
C8A—C13A—C14A	122.81 (9)	C5C—C4C—C3C	119.55 (10)
C12A—C13A—C14A	119.69 (9)	C5C—C4C—H4CA	120.2
C13A—C14A—C1A	112.91 (8)	C3C—C4C—H4CA	120.2
C13A—C14A—C16A	109.53 (8)	C4C—C5C—C6C	120.42 (10)
C1A—C14A—C16A	109.86 (9)	C4C—C5C—H5CA	119.8
C13A—C14A—C15A	107.85 (8)	C6C—C5C—H5CA	119.8
C1A—C14A—C15A	107.88 (8)	C1C—C6C—C5C	120.78 (10)
C16A—C14A—C15A	108.70 (9)	C1C—C6C—C7C	121.40 (9)
C14A—C15A—H15A	109.5	C5C—C6C—C7C	117.82 (9)
C14A—C15A—H15B	109.5	O1C—C7C—C8C	120.98 (10)
H15A—C15A—H15B	109.5	O1C—C7C—C6C	121.04 (10)
C14A—C15A—H15C	109.5	C8C—C7C—C6C	117.98 (9)
H15A—C15A—H15C	109.5	C13C—C8C—C9C	120.82 (9)
H15B—C15A—H15C	109.5	C13C—C8C—C7C	121.05 (9)
C14A—C16A—H16A	109.5	C9C—C8C—C7C	118.12 (9)

C14A—C16A—H16B	109.5	C10C—C9C—C8C	120.96 (10)
H16A—C16A—H16B	109.5	C10C—C9C—H9CA	119.5
C14A—C16A—H16C	109.5	C8C—C9C—H9CA	119.5
H16A—C16A—H16C	109.5	C9C—C10C—C11C	118.83 (10)
H16B—C16A—H16C	109.5	C9C—C10C—H10C	120.6
C6B—C1B—C2B	117.54 (9)	C11C—C10C—H10C	120.6
C6B—C1B—C14B	122.94 (9)	C12C—C11C—C10C	120.45 (10)
C2B—C1B—C14B	119.46 (9)	C12C—C11C—H11C	119.8
C3B—C2B—C1B	121.51 (10)	C10C—C11C—H11C	119.8
C3B—C2B—H2BA	119.2	C11C—C12C—C13C	121.79 (10)
C1B—C2B—H2BA	119.2	C11C—C12C—H12C	119.1
C4B—C3B—C2B	120.13 (10)	C13C—C12C—H12C	119.1
C4B—C3B—H3BA	119.9	C8C—C13C—C12C	117.14 (9)
C2B—C3B—H3BA	119.9	C8C—C13C—C14C	123.28 (9)
C5B—C4B—C3B	119.62 (10)	C12C—C13C—C14C	119.57 (9)
C5B—C4B—H4BA	120.2	C1C—C14C—C13C	113.28 (8)
C3B—C4B—H4BA	120.2	C1C—C14C—C15C	108.61 (8)
C4B—C5B—C6B	120.39 (10)	C13C—C14C—C15C	108.91 (8)
C4B—C5B—H5BA	119.8	C1C—C14C—C16C	108.38 (8)
C6B—C5B—H5BA	119.8	C13C—C14C—C16C	108.09 (8)
C1B—C6B—C5B	120.78 (10)	C15C—C14C—C16C	109.54 (8)
C1B—C6B—C7B	121.12 (9)	C14C—C15C—H15G	109.5
C5B—C6B—C7B	118.08 (9)	C14C—C15C—H15H	109.5
O1B—C7B—C6B	120.70 (10)	H15G—C15C—H15H	109.5
O1B—C7B—C8B	121.46 (10)	C14C—C15C—H15I	109.5
C6B—C7B—C8B	117.83 (9)	H15G—C15C—H15I	109.5
C13B—C8B—C9B	120.44 (10)	H15H—C15C—H15I	109.5
C13B—C8B—C7B	121.31 (9)	C14C—C16C—H16G	109.5
C9B—C8B—C7B	118.25 (9)	C14C—C16C—H16H	109.5
C10B—C9B—C8B	120.80 (10)	H16G—C16C—H16H	109.5
C10B—C9B—H9BA	119.6	C14C—C16C—H16I	109.5
C8B—C9B—H9BA	119.6	H16G—C16C—H16I	109.5
C9B—C10B—C11B	119.22 (10)	H16H—C16C—H16I	109.5
C9B—C10B—H10B	120.4		
C6A—C1A—C2A—C3A	-0.26 (15)	C8B—C9B—C10B—C11B	0.04 (16)
C14A—C1A—C2A—C3A	-177.26 (9)	C9B—C10B—C11B—C12B	-0.10 (16)
C1A—C2A—C3A—C4A	1.80 (16)	C10B—C11B—C12B—C13B	-0.32 (16)
C2A—C3A—C4A—C5A	-1.52 (16)	C9B—C8B—C13B—C12B	-0.83 (15)
C3A—C4A—C5A—C6A	-0.26 (16)	C7B—C8B—C13B—C12B	178.22 (9)
C2A—C1A—C6A—C5A	-1.52 (14)	C9B—C8B—C13B—C14B	-178.93 (9)
C14A—C1A—C6A—C5A	175.40 (9)	C7B—C8B—C13B—C14B	0.12 (15)
C2A—C1A—C6A—C7A	176.71 (9)	C11B—C12B—C13B—C8B	0.77 (15)
C14A—C1A—C6A—C7A	-6.38 (14)	C11B—C12B—C13B—C14B	178.93 (10)
C4A—C5A—C6A—C1A	1.81 (15)	C6B—C1B—C14B—C13B	10.95 (14)
C4A—C5A—C6A—C7A	-176.48 (10)	C2B—C1B—C14B—C13B	-171.94 (9)
C1A—C6A—C7A—O1A	175.70 (10)	C6B—C1B—C14B—C15B	133.38 (10)
C5A—C6A—C7A—O1A	-6.02 (15)	C2B—C1B—C14B—C15B	-49.51 (12)

C1A—C6A—C7A—C8A	-4.97 (14)	C6B—C1B—C14B—C16B	-108.02 (11)
C5A—C6A—C7A—C8A	173.31 (9)	C2B—C1B—C14B—C16B	69.10 (12)
O1A—C7A—C8A—C13A	-172.58 (10)	C8B—C13B—C14B—C1B	-8.51 (14)
C6A—C7A—C8A—C13A	8.10 (14)	C12B—C13B—C14B—C1B	173.42 (9)
O1A—C7A—C8A—C9A	6.77 (15)	C8B—C13B—C14B—C15B	-131.14 (10)
C6A—C7A—C8A—C9A	-172.56 (9)	C12B—C13B—C14B—C15B	50.80 (12)
C13A—C8A—C9A—C10A	0.37 (15)	C8B—C13B—C14B—C16B	110.33 (11)
C7A—C8A—C9A—C10A	-178.98 (9)	C12B—C13B—C14B—C16B	-67.73 (12)
C8A—C9A—C10A—C11A	1.02 (16)	C6C—C1C—C2C—C3C	1.30 (16)
C9A—C10A—C11A—C12A	-0.68 (16)	C14C—C1C—C2C—C3C	-178.80 (10)
C10A—C11A—C12A—C13A	-1.05 (17)	C1C—C2C—C3C—C4C	-0.35 (17)
C9A—C8A—C13A—C12A	-2.02 (15)	C2C—C3C—C4C—C5C	-0.68 (18)
C7A—C8A—C13A—C12A	177.31 (9)	C3C—C4C—C5C—C6C	0.72 (18)
C9A—C8A—C13A—C14A	-179.17 (9)	C2C—C1C—C6C—C5C	-1.25 (15)
C7A—C8A—C13A—C14A	0.16 (15)	C14C—C1C—C6C—C5C	178.85 (9)
C11A—C12A—C13A—C8A	2.37 (16)	C2C—C1C—C6C—C7C	179.17 (9)
C11A—C12A—C13A—C14A	179.62 (10)	C14C—C1C—C6C—C7C	-0.73 (15)
C8A—C13A—C14A—C1A	-10.51 (13)	C4C—C5C—C6C—C1C	0.27 (16)
C12A—C13A—C14A—C1A	172.40 (9)	C4C—C5C—C6C—C7C	179.86 (10)
C8A—C13A—C14A—C16A	-133.28 (10)	C1C—C6C—C7C—O1C	-178.21 (10)
C12A—C13A—C14A—C16A	49.63 (12)	C5C—C6C—C7C—O1C	2.20 (16)
C8A—C13A—C14A—C15A	108.59 (11)	C1C—C6C—C7C—C8C	2.14 (14)
C12A—C13A—C14A—C15A	-68.50 (12)	C5C—C6C—C7C—C8C	-177.45 (9)
C2A—C1A—C14A—C13A	-169.52 (9)	O1C—C7C—C8C—C13C	179.49 (10)
C6A—C1A—C14A—C13A	13.65 (13)	C6C—C7C—C8C—C13C	-0.86 (14)
C2A—C1A—C14A—C16A	-46.94 (12)	O1C—C7C—C8C—C9C	-1.52 (15)
C6A—C1A—C14A—C16A	136.23 (10)	C6C—C7C—C8C—C9C	178.13 (9)
C2A—C1A—C14A—C15A	71.40 (12)	C13C—C8C—C9C—C10C	0.14 (16)
C6A—C1A—C14A—C15A	-105.43 (11)	C7C—C8C—C9C—C10C	-178.85 (10)
C6B—C1B—C2B—C3B	-0.66 (16)	C8C—C9C—C10C—C11C	0.35 (16)
C14B—C1B—C2B—C3B	-177.93 (10)	C9C—C10C—C11C—C12C	-0.69 (17)
C1B—C2B—C3B—C4B	1.40 (17)	C10C—C11C—C12C—C13C	0.55 (17)
C2B—C3B—C4B—C5B	-0.92 (17)	C9C—C8C—C13C—C12C	-0.28 (14)
C3B—C4B—C5B—C6B	-0.26 (17)	C7C—C8C—C13C—C12C	178.68 (9)
C2B—C1B—C6B—C5B	-0.53 (15)	C9C—C8C—C13C—C14C	179.19 (9)
C14B—C1B—C6B—C5B	176.64 (9)	C7C—C8C—C13C—C14C	-1.85 (15)
C2B—C1B—C6B—C7B	177.92 (9)	C11C—C12C—C13C—C8C	-0.06 (15)
C14B—C1B—C6B—C7B	-4.92 (15)	C11C—C12C—C13C—C14C	-179.55 (10)
C4B—C5B—C6B—C1B	1.00 (16)	C6C—C1C—C14C—C13C	-1.78 (13)
C4B—C5B—C6B—C7B	-177.50 (10)	C2C—C1C—C14C—C13C	178.33 (9)
C1B—C6B—C7B—O1B	176.03 (10)	C6C—C1C—C14C—C15C	-122.91 (10)
C5B—C6B—C7B—O1B	-5.48 (15)	C2C—C1C—C14C—C15C	57.19 (12)
C1B—C6B—C7B—C8B	-4.30 (15)	C6C—C1C—C14C—C16C	118.17 (10)
C5B—C6B—C7B—C8B	174.19 (9)	C2C—C1C—C14C—C16C	-61.73 (12)
O1B—C7B—C8B—C13B	-173.65 (10)	C8C—C13C—C14C—C1C	3.08 (13)
C6B—C7B—C8B—C13B	6.68 (14)	C12C—C13C—C14C—C1C	-177.46 (9)
O1B—C7B—C8B—C9B	5.42 (15)	C8C—C13C—C14C—C15C	124.05 (10)
C6B—C7B—C8B—C9B	-174.25 (9)	C12C—C13C—C14C—C15C	-56.49 (12)

C13B—C8B—C9B—C10B	0.44 (16)	C8C—C13C—C14C—C16C	-117.03 (10)
C7B—C8B—C9B—C10B	-178.64 (10)	C12C—C13C—C14C—C16C	62.43 (12)

Hydrogen-bond geometry (Å, °)

Cg1, Cg7, Cg13 and Cg15 are the centroids of the C1A–C6A, C1B–C6B, C1C–C6C and C8C–C13C rings, respectively.

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
C10 <i>A</i> —H10 <i>A</i> ...O1 <i>C</i> ⁱ	0.93	2.52	3.4097 (16)	160
C10 <i>C</i> —H10 <i>C</i> ...O1 <i>B</i> ⁱⁱ	0.93	2.59	3.4374 (18)	151
C3 <i>C</i> —H3 <i>CA</i> ...O1 <i>A</i> ⁱⁱⁱ	0.93	2.45	3.2287 (17)	141
C15 <i>B</i> —H15 <i>D</i> ...O1 <i>B</i> ⁱⁱ	0.96	2.43	3.3568 (18)	161
C16 <i>A</i> —H16 <i>A</i> ...O1 <i>A</i> ^{iv}	0.96	2.48	3.3934 (16)	159
C3 <i>A</i> —H3 <i>AA</i> ...Cg13 ^v	0.93	2.69	3.3996 (14)	134
C3 <i>B</i> —H3 <i>BA</i> ...Cg13 ^{vi}	0.93	2.77	3.3715 (14)	123
C15 <i>A</i> —H15 <i>B</i> ...Cg15 ^v	0.96	2.71	3.4915 (15)	139
C15 <i>C</i> —H15 <i>G</i> ...Cg1 ^{vii}	0.96	2.96	3.8656 (13)	158
C16 <i>C</i> —H16 <i>G</i> ...Cg7 ^{vi}	0.96	2.91	3.7432 (14)	145

Symmetry codes: (i) $x-1, y, z$; (ii) $-x+1, -y+1, -z$; (iii) $-x+2, -y+1, -z+1$; (iv) $-x+1, -y+2, -z+1$; (v) $x, y+1, z$; (vi) $-x+2, -y+1, -z$; (vii) $x, y-1, z$.