

10-(4-Methylbenzylidene)anthracen-9(10H)-one

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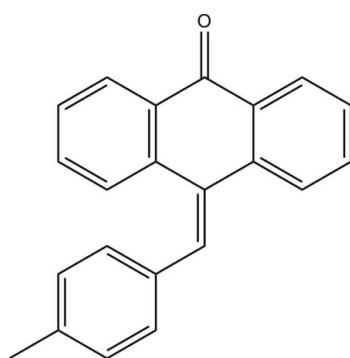
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Key indicators: single-crystal X-ray study; $T = 100$ K; mean $\sigma(C-C) = 0.002$ Å; R factor = 0.043; wR factor = 0.122; data-to-parameter ratio = 13.2.

In the title compound, $C_{22}H_{16}O$, the six-membered ring within the anthrone moiety adopts a shallow boat conformation, with puckering parameters $Q = 0.2860$ (17) Å, $\Theta = 99.1$ (3)° and $\Phi = 114.8$ (3)°. The dihedral angle between the outer benzene rings is 26.53 (8)°. The mean plane through the anthrone ring system makes a dihedral angle of 38.73 (6)° with the pendant benzene ring. In the crystal, molecules are linked by C–H···O hydrogen bonds into zigzag chains propagating along the *c*-axis direction and weak C–H···π interactions further consolidate the structure.

Related literature

For a related structure and background to anthrone derivatives, see: Arumugam *et al.* (2011). For related structures, see: Wen & Li (2008); Zhou *et al.* (2004). For the synthesis, see: Prinz *et al.* (2003). For ring conformations, see: Cremer & Pople (1975). For the stability of the temperature controller used in the data collection, see: Cosier & Glazer (1986).



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Experimental

Crystal data

$C_{22}H_{16}O$	$V = 1554.43$ (4) Å ³
$M_r = 296.35$	$Z = 4$
Orthorhombic, $Pna2_1$	Mo $K\alpha$ radiation
$a = 7.2959$ (1) Å	$\mu = 0.08$ mm ⁻¹
$b = 16.3853$ (2) Å	$T = 100$ K
$c = 13.0028$ (2) Å	$0.33 \times 0.27 \times 0.21$ mm

Data collection

Bruker SMART APEXII CCD diffractometer	20184 measured reflections
Absorption correction: multi-scan (<i>SADABS</i> ; Bruker, 2009)	2757 independent reflections
$(SADABS$; Bruker, 2009)	2575 reflections with $I > 2\sigma(I)$
$T_{\min} = 0.975$, $T_{\max} = 0.985$	$R_{\text{int}} = 0.030$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.043$	H-atom parameters constrained
$wR(F^2) = 0.122$	$\Delta\rho_{\max} = 0.40$ e Å ⁻³
$S = 1.09$	$\Delta\rho_{\min} = -0.19$ e Å ⁻³
2757 reflections	Absolute structure: Flack (1983), 2319 Friedel pairs
209 parameters	Flack parameter: 0 (10)
1 restraint	

Table 1

Hydrogen-bond geometry (Å, °).

Cg1 and *Cg2* are the centroids of the C1–C6 and C16–C21 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>
C3–H3A···O1 ⁱ	0.95	2.35	3.275 (2)	164
C22–H22C··· <i>Cg1</i> ⁱⁱ	0.98	2.94	3.726 (2)	138
C17–H17A··· <i>Cg2</i> ⁱⁱⁱ	0.95	2.76	3.5073 (16)	136

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

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: HB6592).

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supporting information

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S1. Comment

As part of our ongoing studies of anthrone derivatives (Arumugam *et al.*, 2011), we have undertaken the X-ray crystal structure determination of the title compound, (I).

In the molecular structure (Fig 1), the six-membered ring (C1/C6–C8/C13/C14) within the anthracene moiety adopts a boat conformation with puckering parameters $Q=0.2860$ (17) Å, $\Theta=99.1$ (3)° and $\Phi=114.8$ (3)° (Cremer & Pople, 1975). This differs from the planar anthracene ring reported in related structure (Arumugam *et al.*, 2011). The mean plane through the anthracene ring (C1–C14) makes a dihedral angle of 38.73 (6)° with the phenyl ring (C16–C21). The bond lengths and angles are comparable those in the related structure (Wen & Li, 2008; Zhou *et al.*, 2004).

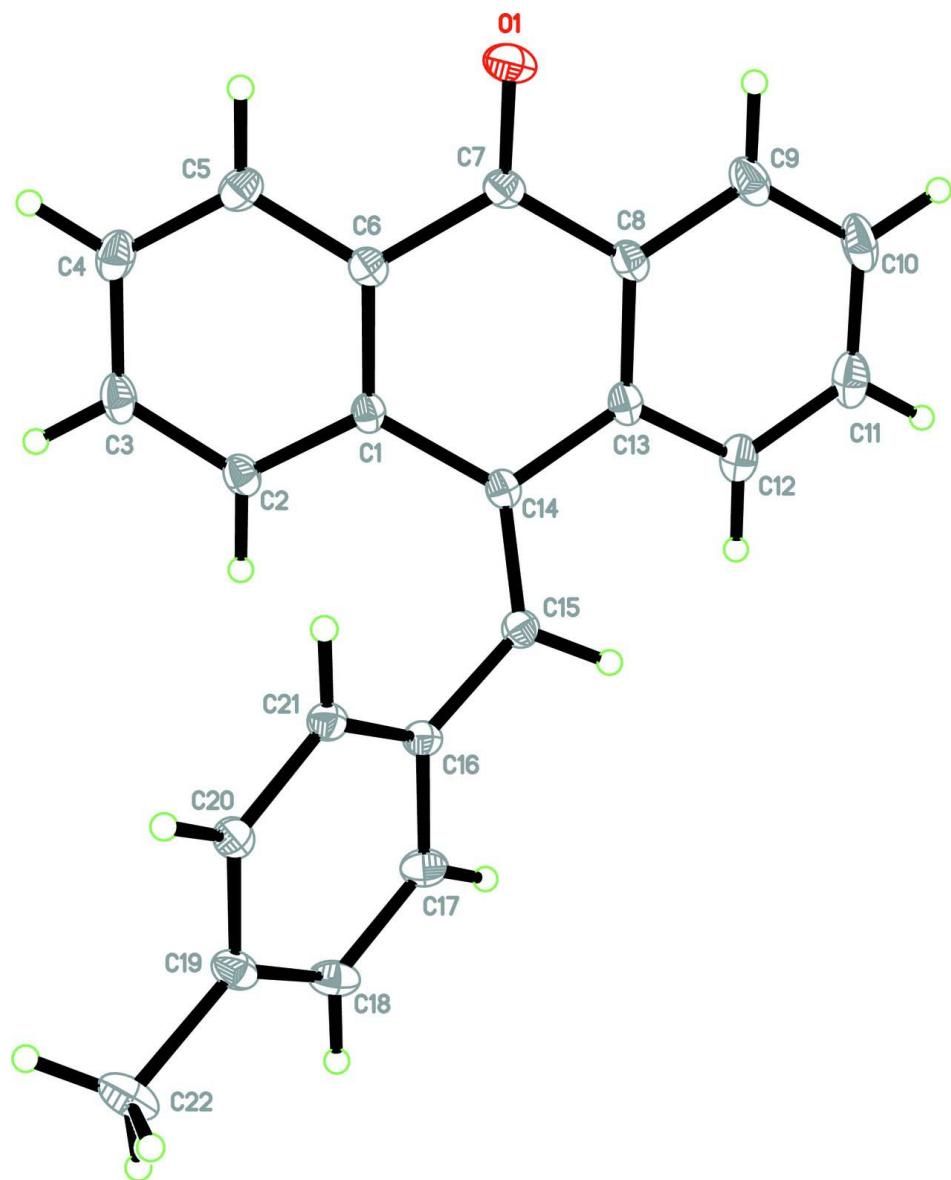
The crystal packing is shown in Fig. 2. The molecules are linked by the intermolecular C3—H3A···O1 hydrogen bonds (Table 1) into infinite one dimensional zigzag chain along the *c*-axis. In addition, the crystal structure are further stabilized by the intermolecular C22—H22C···*Cg1* and C17—H17A···*Cg2* (Table 1) interactions (*Cg1* and *Cg2* are the centroids of C1–C6 and C16–C21 rings, respectively).

S2. Experimental

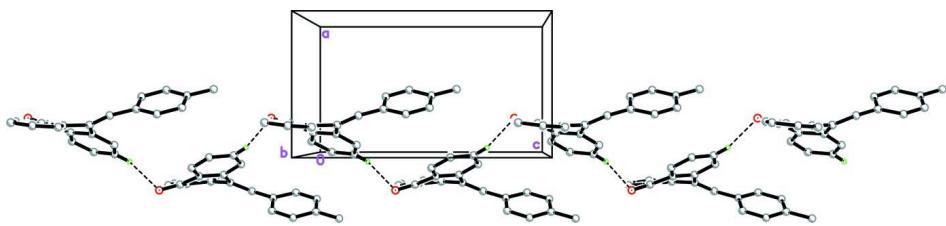
The title compound was synthesized as reported (Prinz *et al.*, 2003) and crystallized EtOAc by slow evaporation technique.

S3. Refinement

All H atoms were positioned geometrically [C–H = 0.95 and 0.98 Å] and refined using a riding model with $U_{\text{iso}}(\text{H}) = 1.2$ and 1.5 $U_{\text{eq}}(\text{C})$. A rotating group model was applied to the methyl groups. A total of 2319 Freidel pairs were used to determine the absolute structure.

**Figure 1**

The molecular structure of the title compound, showing 30% probability displacement ellipsoids.

**Figure 2**

The crystal packing of the title compound. The H atoms not involved in the intermolecular interactions (dashed lines) have been omitted for clarity.

10-(4-Methylbenzylidene)anthracen-9(10*H*)-one*Crystal data*

$C_{22}H_{16}O$
 $M_r = 296.35$
Orthorhombic, $Pna2_1$
Hall symbol: P 2c -2n
 $a = 7.2959$ (1) Å
 $b = 16.3853$ (2) Å
 $c = 13.0028$ (2) Å
 $V = 1554.43$ (4) Å³
 $Z = 4$

$F(000) = 624$
 $D_x = 1.266 \text{ Mg m}^{-3}$
Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
Cell parameters from 8371 reflections
 $\theta = 3.1\text{--}31.7^\circ$
 $\mu = 0.08 \text{ mm}^{-1}$
 $T = 100$ K
Block, yellow
 $0.33 \times 0.27 \times 0.21$ mm

Data collection

Bruker SMART APEXII CCD
diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
 φ and ω scans
Absorption correction: multi-scan
(SADABS; Bruker, 2009)
 $T_{\min} = 0.975$, $T_{\max} = 0.985$

20184 measured reflections
2757 independent reflections
2575 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.030$
 $\theta_{\max} = 31.8^\circ$, $\theta_{\min} = 2.0^\circ$
 $h = -10 \rightarrow 10$
 $k = -24 \rightarrow 21$
 $l = -19 \rightarrow 17$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.043$
 $wR(F^2) = 0.122$
 $S = 1.09$
2757 reflections
209 parameters
1 restraint
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
 $w = 1/[\sigma^2(F_o^2) + (0.0831P)^2 + 0.1277P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.002$
 $\Delta\rho_{\max} = 0.40 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.19 \text{ e } \text{\AA}^{-3}$
Absolute structure: Flack (1983), 2319 Friedel
pairs
Absolute structure parameter: 0 (10)

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 (Å²)

	x	y	z	$U_{\text{iso}}^* / U_{\text{eq}}$
O1	0.22961 (18)	0.09172 (9)	0.36336 (11)	0.0332 (3)

C1	0.34392 (19)	0.11506 (9)	0.63231 (12)	0.0204 (3)
C2	0.4267 (2)	0.07120 (10)	0.71248 (13)	0.0249 (3)
H2A	0.4357	0.0949	0.7789	0.030*
C3	0.4956 (2)	-0.00689 (11)	0.69531 (15)	0.0297 (4)
H3A	0.5527	-0.0357	0.7500	0.036*
C4	0.4818 (2)	-0.04321 (11)	0.59886 (16)	0.0311 (4)
H4A	0.5270	-0.0969	0.5880	0.037*
C5	0.4016 (2)	-0.00049 (10)	0.51897 (15)	0.0274 (3)
H5A	0.3917	-0.0250	0.4530	0.033*
C6	0.3352 (2)	0.07847 (10)	0.53448 (12)	0.0223 (3)
C7	0.2696 (2)	0.12531 (10)	0.44437 (12)	0.0234 (3)
C8	0.2672 (2)	0.21516 (11)	0.45515 (12)	0.0239 (3)
C9	0.2547 (3)	0.26344 (13)	0.36613 (16)	0.0328 (4)
H9A	0.2415	0.2381	0.3008	0.039*
C10	0.2613 (3)	0.34737 (13)	0.37306 (18)	0.0384 (5)
H10A	0.2514	0.3799	0.3129	0.046*
C11	0.2828 (2)	0.38405 (12)	0.46872 (18)	0.0351 (4)
H11A	0.2905	0.4418	0.4733	0.042*
C12	0.2931 (2)	0.33765 (10)	0.55740 (16)	0.0281 (3)
H12A	0.3071	0.3638	0.6221	0.034*
C13	0.2829 (2)	0.25202 (10)	0.55238 (13)	0.0227 (3)
C14	0.2792 (2)	0.20019 (9)	0.64518 (12)	0.0203 (3)
C15	0.2111 (2)	0.23176 (9)	0.73396 (12)	0.0223 (3)
H15A	0.1904	0.2890	0.7325	0.027*
C16	0.1643 (2)	0.19219 (9)	0.83166 (12)	0.0220 (3)
C17	0.2006 (2)	0.23303 (10)	0.92404 (14)	0.0274 (3)
H17A	0.2577	0.2851	0.9224	0.033*
C18	0.1542 (2)	0.19832 (12)	1.01808 (13)	0.0293 (3)
H18A	0.1828	0.2265	1.0799	0.035*
C19	0.0662 (2)	0.12249 (11)	1.02327 (13)	0.0269 (3)
C20	0.0247 (2)	0.08341 (10)	0.93057 (13)	0.0240 (3)
H20A	-0.0385	0.0327	0.9321	0.029*
C21	0.0736 (2)	0.11680 (9)	0.83647 (12)	0.0223 (3)
H21A	0.0454	0.0884	0.7748	0.027*
C22	0.0189 (3)	0.08462 (15)	1.12512 (15)	0.0386 (4)
H22A	0.0738	0.1168	1.1807	0.058*
H22B	0.0665	0.0287	1.1276	0.058*
H22C	-0.1146	0.0836	1.1335	0.058*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0306 (6)	0.0475 (7)	0.0215 (5)	0.0048 (5)	-0.0009 (5)	-0.0047 (5)
C1	0.0160 (6)	0.0216 (6)	0.0235 (6)	0.0018 (4)	0.0013 (5)	0.0038 (5)
C2	0.0206 (6)	0.0296 (7)	0.0245 (6)	0.0027 (6)	-0.0003 (6)	0.0064 (6)
C3	0.0224 (6)	0.0301 (7)	0.0364 (9)	0.0063 (6)	0.0030 (6)	0.0117 (7)
C4	0.0247 (7)	0.0260 (7)	0.0425 (9)	0.0062 (6)	0.0062 (7)	0.0039 (7)
C5	0.0226 (7)	0.0275 (7)	0.0321 (8)	0.0018 (6)	0.0051 (6)	-0.0029 (6)

C6	0.0172 (6)	0.0272 (7)	0.0225 (6)	0.0009 (5)	0.0019 (5)	0.0028 (5)
C7	0.0178 (6)	0.0317 (7)	0.0207 (7)	0.0026 (5)	0.0025 (5)	0.0010 (6)
C8	0.0181 (6)	0.0317 (7)	0.0219 (7)	0.0035 (5)	0.0034 (5)	0.0074 (6)
C9	0.0257 (7)	0.0466 (10)	0.0261 (7)	0.0071 (7)	0.0057 (6)	0.0138 (8)
C10	0.0300 (8)	0.0449 (10)	0.0403 (10)	0.0085 (7)	0.0091 (7)	0.0259 (9)
C11	0.0244 (8)	0.0312 (8)	0.0495 (11)	0.0028 (6)	0.0096 (7)	0.0173 (8)
C12	0.0205 (7)	0.0250 (7)	0.0387 (9)	-0.0004 (5)	0.0062 (6)	0.0080 (7)
C13	0.0164 (6)	0.0252 (7)	0.0264 (7)	0.0012 (5)	0.0019 (5)	0.0071 (6)
C14	0.0168 (6)	0.0221 (6)	0.0221 (6)	-0.0005 (4)	-0.0005 (5)	0.0030 (5)
C15	0.0225 (7)	0.0215 (6)	0.0227 (7)	-0.0007 (5)	0.0008 (5)	0.0006 (5)
C16	0.0219 (7)	0.0242 (6)	0.0198 (6)	0.0007 (5)	0.0010 (6)	-0.0011 (5)
C17	0.0260 (7)	0.0315 (8)	0.0249 (7)	-0.0038 (6)	0.0009 (6)	-0.0060 (6)
C18	0.0264 (8)	0.0416 (9)	0.0198 (7)	-0.0010 (6)	-0.0008 (6)	-0.0062 (6)
C19	0.0205 (7)	0.0394 (8)	0.0208 (6)	0.0018 (6)	0.0013 (6)	0.0031 (6)
C20	0.0208 (6)	0.0272 (7)	0.0239 (7)	0.0019 (5)	0.0011 (6)	0.0025 (6)
C21	0.0224 (6)	0.0235 (6)	0.0210 (6)	0.0001 (5)	0.0020 (5)	-0.0002 (5)
C22	0.0322 (9)	0.0613 (12)	0.0224 (8)	-0.0067 (8)	0.0017 (7)	0.0081 (8)

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

O1—C7	1.224 (2)	C11—H11A	0.9500
C1—C2	1.403 (2)	C12—C13	1.407 (2)
C1—C6	1.408 (2)	C12—H12A	0.9500
C1—C14	1.482 (2)	C13—C14	1.476 (2)
C2—C3	1.393 (2)	C14—C15	1.359 (2)
C2—H2A	0.9500	C15—C16	1.467 (2)
C3—C4	1.392 (3)	C15—H15A	0.9500
C3—H3A	0.9500	C16—C17	1.400 (2)
C4—C5	1.382 (3)	C16—C21	1.403 (2)
C4—H4A	0.9500	C17—C18	1.390 (2)
C5—C6	1.396 (2)	C17—H17A	0.9500
C5—H5A	0.9500	C18—C19	1.400 (3)
C6—C7	1.480 (2)	C18—H18A	0.9500
C7—C8	1.479 (2)	C19—C20	1.398 (2)
C8—C13	1.406 (2)	C19—C22	1.503 (3)
C8—C9	1.405 (2)	C20—C21	1.387 (2)
C9—C10	1.379 (3)	C20—H20A	0.9500
C9—H9A	0.9500	C21—H21A	0.9500
C10—C11	1.390 (4)	C22—H22A	0.9800
C10—H10A	0.9500	C22—H22B	0.9800
C11—C12	1.383 (3)	C22—H22C	0.9800
C2—C1—C6	118.21 (14)	C13—C12—H12A	119.8
C2—C1—C14	122.37 (15)	C8—C13—C12	118.33 (15)
C6—C1—C14	119.25 (14)	C8—C13—C14	119.12 (14)
C3—C2—C1	120.46 (16)	C12—C13—C14	122.48 (15)
C3—C2—H2A	119.8	C15—C14—C13	118.82 (13)
C1—C2—H2A	119.8	C15—C14—C1	124.79 (14)

C4—C3—C2	120.73 (15)	C13—C14—C1	116.33 (14)
C4—C3—H3A	119.6	C14—C15—C16	130.74 (13)
C2—C3—H3A	119.6	C14—C15—H15A	114.6
C5—C4—C3	119.43 (15)	C16—C15—H15A	114.6
C5—C4—H4A	120.3	C17—C16—C21	118.16 (14)
C3—C4—H4A	120.3	C17—C16—C15	119.18 (13)
C4—C5—C6	120.50 (17)	C21—C16—C15	122.54 (14)
C4—C5—H5A	119.8	C18—C17—C16	120.85 (15)
C6—C5—H5A	119.8	C18—C17—H17A	119.6
C5—C6—C1	120.63 (15)	C16—C17—H17A	119.6
C5—C6—C7	118.58 (15)	C17—C18—C19	121.12 (15)
C1—C6—C7	120.60 (14)	C17—C18—H18A	119.4
O1—C7—C8	121.76 (15)	C19—C18—H18A	119.4
O1—C7—C6	121.69 (15)	C20—C19—C18	117.66 (15)
C8—C7—C6	116.42 (14)	C20—C19—C22	121.40 (16)
C13—C8—C9	120.29 (16)	C18—C19—C22	120.94 (16)
C13—C8—C7	120.79 (13)	C21—C20—C19	121.62 (14)
C9—C8—C7	118.89 (16)	C21—C20—H20A	119.2
C10—C9—C8	120.4 (2)	C19—C20—H20A	119.2
C10—C9—H9A	119.8	C20—C21—C16	120.53 (15)
C8—C9—H9A	119.8	C20—C21—H21A	119.7
C9—C10—C11	119.57 (17)	C16—C21—H21A	119.7
C9—C10—H10A	120.2	C19—C22—H22A	109.5
C11—C10—H10A	120.2	C19—C22—H22B	109.5
C12—C11—C10	120.96 (17)	H22A—C22—H22B	109.5
C12—C11—H11A	119.5	C19—C22—H22C	109.5
C10—C11—H11A	119.5	H22A—C22—H22C	109.5
C11—C12—C13	120.45 (18)	H22B—C22—H22C	109.5
C11—C12—H12A	119.8		
C6—C1—C2—C3	-0.9 (2)	C9—C8—C13—C14	174.59 (15)
C14—C1—C2—C3	-176.04 (15)	C7—C8—C13—C14	-7.4 (2)
C1—C2—C3—C4	-0.8 (3)	C11—C12—C13—C8	1.7 (2)
C2—C3—C4—C5	1.2 (3)	C11—C12—C13—C14	-175.30 (15)
C3—C4—C5—C6	0.1 (3)	C8—C13—C14—C15	-150.02 (15)
C4—C5—C6—C1	-1.8 (2)	C12—C13—C14—C15	27.0 (2)
C4—C5—C6—C7	173.23 (15)	C8—C13—C14—C1	27.30 (19)
C2—C1—C6—C5	2.2 (2)	C12—C13—C14—C1	-155.71 (14)
C14—C1—C6—C5	177.49 (14)	C2—C1—C14—C15	-32.6 (2)
C2—C1—C6—C7	-172.75 (14)	C6—C1—C14—C15	152.29 (16)
C14—C1—C6—C7	2.6 (2)	C2—C1—C14—C13	150.25 (14)
C5—C6—C7—O1	18.3 (2)	C6—C1—C14—C13	-24.9 (2)
C1—C6—C7—O1	-166.66 (15)	C13—C14—C15—C16	169.27 (16)
C5—C6—C7—C8	-157.66 (14)	C1—C14—C15—C16	-7.8 (3)
C1—C6—C7—C8	17.4 (2)	C14—C15—C16—C17	142.24 (19)
O1—C7—C8—C13	169.11 (15)	C14—C15—C16—C21	-41.8 (3)
C6—C7—C8—C13	-14.9 (2)	C21—C16—C17—C18	2.4 (2)
O1—C7—C8—C9	-12.8 (2)	C15—C16—C17—C18	178.48 (17)

C6—C7—C8—C9	163.10 (15)	C16—C17—C18—C19	-1.5 (3)
C13—C8—C9—C10	1.3 (3)	C17—C18—C19—C20	-0.7 (3)
C7—C8—C9—C10	-176.73 (16)	C17—C18—C19—C22	179.07 (18)
C8—C9—C10—C11	0.8 (3)	C18—C19—C20—C21	2.0 (2)
C9—C10—C11—C12	-1.6 (3)	C22—C19—C20—C21	-177.81 (17)
C10—C11—C12—C13	0.3 (3)	C19—C20—C21—C16	-1.1 (2)
C9—C8—C13—C12	-2.5 (2)	C17—C16—C21—C20	-1.1 (2)
C7—C8—C13—C12	175.48 (14)	C15—C16—C21—C20	-177.09 (14)

Hydrogen-bond geometry (Å, °)

Cg1 and Cg2 are the centroids of the C1—C6 and C16—C21 rings, respectively.

D—H···A	D—H	H···A	D···A	D—H···A
C3—H3A···O1 ⁱ	0.95	2.35	3.275 (2)	164
C22—H22C···Cg1 ⁱⁱ	0.98	2.94	3.726 (2)	138
C17—H17A···Cg2 ⁱⁱⁱ	0.95	2.76	3.5073 (16)	136

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