

Ethyl 4-(4-methoxyphenyl)-2-oxo-6-phenylcyclohex-3-ene-1-carboxylate

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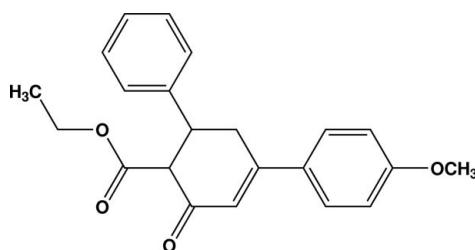
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Key indicators: single-crystal X-ray study; $T = 100\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.002\text{ \AA}$; disorder in main residue; R factor = 0.058; wR factor = 0.169; data-to-parameter ratio = 24.9.

The asymmetric unit of the title compound, $C_{22}H_{22}O_4$, consists of two independent molecules (A and B) which differ significantly in the orientations of ethyl carboxylate groups. The phenyl ring in molecule B is disordered over two orientations with occupancies of 0.55 (2) and 0.45 (2). The cyclohexenone ring of both molecules adopts an envelope conformation. The dihedral angle between the two aromatic rings is 81.12 (7) $^\circ$ in molecule A and 70.8 (3) $^\circ$ in molecule B [57.5 (4) $^\circ$ in the minor disorder component]. The crystal structure is stabilized by weak intermolecular C–H \cdots O hydrogen bonds and C–H \cdots π interactions.

Related literature

For general background, see: Kalluraya & Rai (2003); Kalluraya & Rahiman (2003). For bond-length data, see: Allen *et al.* (1987). For ring puckering analysis, see: Cremer & Pople (1975).



Experimental

Crystal data

$C_{22}H_{22}O_4$

$M_r = 350.40$

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Triclinic, $P\bar{1}$	$V = 1831.00 (7)\text{ \AA}^3$
$a = 9.6095 (2)\text{ \AA}$	$Z = 4$
$b = 12.5386 (3)\text{ \AA}$	Mo $K\alpha$ radiation
$c = 15.9704 (3)\text{ \AA}$	$\mu = 0.09\text{ mm}^{-1}$
$\alpha = 75.916 (1)^\circ$	$T = 100.0 (1)\text{ K}$
$\beta = 78.879 (1)^\circ$	$0.50 \times 0.28 \times 0.24\text{ mm}$
$\gamma = 88.413 (1)^\circ$	

Data collection

Bruker SMART APEXII CCD area-detector diffractometer	57259 measured reflections
Absorption correction: multi-scan (<i>SADABS</i> ; Bruker, 2005)	13169 independent reflections
$T_{\min} = 0.958$, $T_{\max} = 0.980$	9552 reflections with $I > 2\sigma(I)$
	$R_{\text{int}} = 0.032$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.058$	144 restraints
$wR(F^2) = 0.169$	H-atom parameters constrained
$S = 1.02$	$\Delta\rho_{\text{max}} = 0.92\text{ e \AA}^{-3}$
13169 reflections	$\Delta\rho_{\text{min}} = -0.28\text{ e \AA}^{-3}$
528 parameters	

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

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
C3A–H3A \cdots O2B ⁱ	1.00	2.57	3.4093 (18)	142
C5A–H5AA \cdots O4B ⁱⁱ	0.95	2.55	3.3630 (17)	143
C8A–H8AA \cdots O4B ⁱⁱ	0.95	2.31	3.2446 (18)	166
C14B–H14B \cdots O2A \bar{i}	0.95	2.58	3.500 (5)	163
C16A–H16A \cdots O2B ⁱⁱⁱ	0.95	2.47	3.2789 (18)	143
C17B–H17B \cdots O3A ^{iv}	0.95	2.47	3.116 (9)	125
C21B–H21D \cdots Cg1 ^v	0.98	2.85	3.811 (2)	167
C22B–H22D \cdots Cg2 ^{vi}	0.98	2.87	3.644 (4)	137

Symmetry codes: (i) $-x, -y + 1, -z + 2$; (ii) $-x + 1, -y + 1, -z + 2$; (iii) $x, y + 1, z - 1$; (iv) $x, y - 1, z$; (v) $x + 1, y, z$; (vi) $-x, -y, -z + 2$. Cg1 and Cg2 are the centroids of the C13A–C18A and C13B–C18B rings, respectively.

Data collection: *APEX2* (Bruker, 2005); cell refinement: *SAINT* (Bruker, 2005); 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, 2003).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: CI2722).

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supporting information

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Ethyl 4-(4-methoxyphenyl)-2-oxo-6-phenylcyclohex-3-ene-1-carboxylate

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S1. Comment

Cyclohexenone is an organic compound which is a versatile intermediate used in the synthesis of a variety of chemical products such as pharmaceuticals and fragrances. The Robinson annulation is an organic reaction used to create a cyclic six-membered, β -unsaturated ketones. Different methods including solvent-free synthesis of these compounds have been reported (Kalluraya & Rai, 2003). Cyclohexenone and their derivatives are known for anti-inflammatory and analgesic activities (Kalluraya & Rahiman, 2003).

The asymmetric unit of the title compound consists of two independent molecules (Fig. 1). Bond lengths in the molecules (Fig. 1) are found to have normal values (Allen *et al.*, 1987). The two independent molecules differ significantly in the orientations of ethyl carboxylate groups [$\text{C}4\text{A}$ — $\text{C}3\text{A}$ — $\text{C}19\text{A}$ — $\text{O}1\text{A}$ = 63.32 (14) $^\circ$ and $\text{C}4\text{B}$ — $\text{C}3\text{B}$ — $\text{C}19\text{B}$ — $\text{O}1\text{B}$ = 105.25 (13) $^\circ$]. The phenyl ring in molecule B is disordered over two orientations. The cyclohexenone ring in both molecules adopts an envelope conformation, with puckering parameters Q = 0.494 (1) Å, θ = 52.1 (2) $^\circ$ and φ = 67.4 (2) $^\circ$ for molecule A, and Q = 0.429 (2) Å, θ = 130.1 (2) $^\circ$ and φ = 250.4 (3) $^\circ$ in molecule B (Cremer & Pople, 1975). The dihedral angle between the two aromatic rings is 81.12 (7) $^\circ$ in molecule A and 70.8 (3) $^\circ$ in molecule B [57.5 (4) $^\circ$ in the minor disorder component].

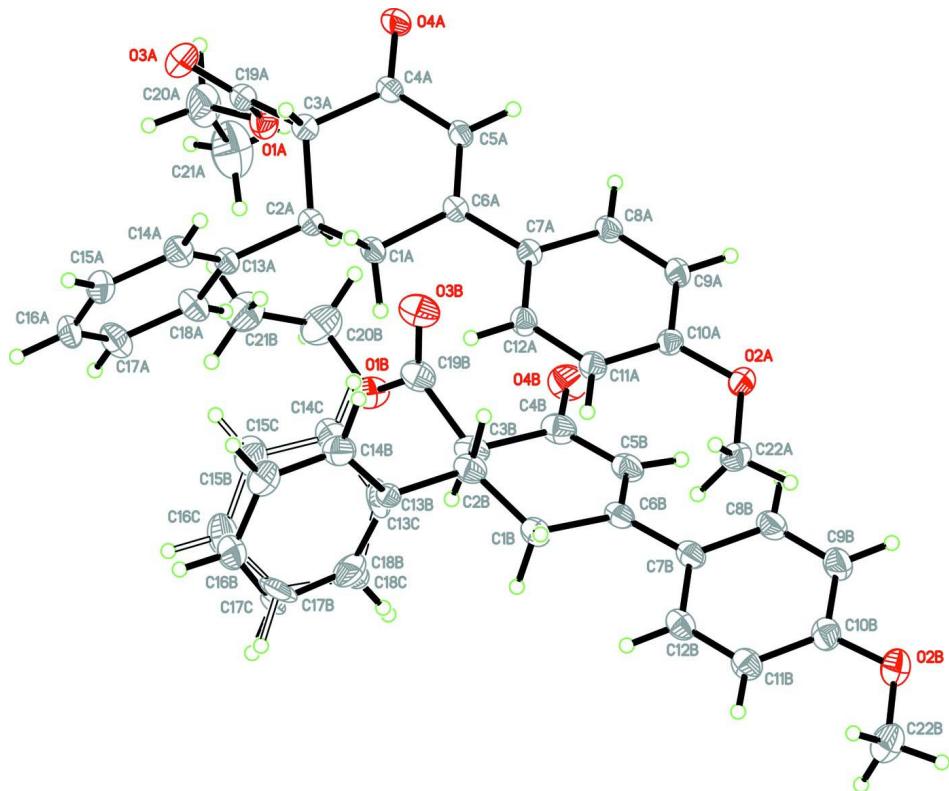
The crystal structure is stabilized by intermolecular weak C—H \cdots O hydrogen bonds (Table 1) and C—H \cdots π interactions (Fig 2).

S2. Experimental

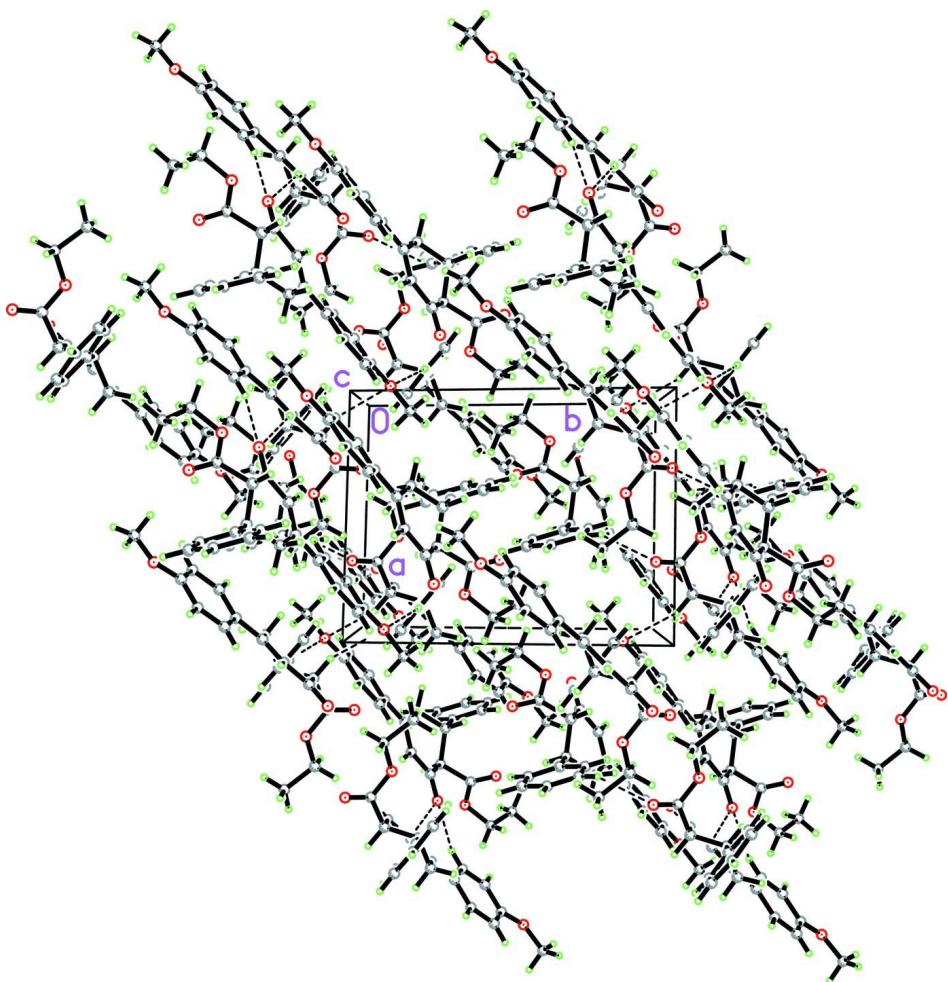
To a solution of 1-phenyl-3-anisyl-prop-2-en-1-one (0.01 mol) in dry acetone (25 ml), dry potassium carbonate (0.04 mol) and ethyl acetoacetate (0.02 mol) in dry acetone (25 ml) were added. The mixture was stirred at room temperature for overnight and was filtered. The solvent from the filtrate on evaporation gave a solid which was recrystallized from a mixture of ethanol-dioxane.

S3. Refinement

The phenyl ring in molecule B is disordered over two orientations with refined occupancies of 0.545 (17) and 0.455 (17). The C-C bond lengths involving the disordered atoms were restrained to be equal and also the U_{ij}^{ij} components of the disordered atoms were approximated to isotropic behaviour. The two orientations were restrained to be planar. H atoms were positioned geometrically ($\text{C}-\text{H}$ = 0.95–1.00 Å) and refined using a riding model, with $U_{iso}(\text{H})$ = 1.2 $U_{eq}(\text{C})$ and 1.5 $U_{eq}(\text{C}_{\text{methyl}})$. A rotating group model was used for the methyl groups. The highest four difference peaks were observed at 0.92, 0.83, 0.87 and 84 Å, respectively, from atoms $\text{C}3\text{B}$, $\text{C}2\text{B}$, $\text{C}2\text{A}$ and $\text{C}3\text{A}$. No suitable disorder model involving these atoms were found.

**Figure 1**

The asymmetric unit of the title compound, showing 50% probability displacement ellipsoids and the atom-numbering scheme. Both disorder component of an independent molecule are shown.

**Figure 2**

The crystal packing of the title compound, viewed down the *c* axis. Only the major disorder component of an independent molecule is shown.

Ethyl 4-(4-methoxyphenyl)-2-oxo-6-phenylcyclohex-3-ene-1-carboxylate

Crystal data

$C_{22}H_{22}O_4$
 $M_r = 350.40$
Triclinic, $P\bar{1}$
Hall symbol: -P 1
 $a = 9.6095 (2) \text{ \AA}$
 $b = 12.5386 (3) \text{ \AA}$
 $c = 15.9704 (3) \text{ \AA}$
 $\alpha = 75.916 (1)^\circ$
 $\beta = 78.879 (1)^\circ$
 $\gamma = 88.413 (1)^\circ$
 $V = 1831.00 (7) \text{ \AA}^3$

$Z = 4$
 $F(000) = 744$
 $D_x = 1.271 \text{ Mg m}^{-3}$
Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$
Cell parameters from 9088 reflections
 $\theta = 2.3\text{--}33.8^\circ$
 $\mu = 0.09 \text{ mm}^{-1}$
 $T = 100 \text{ K}$
Block, colourless
 $0.50 \times 0.28 \times 0.24 \text{ mm}$

Data collection

Bruker SMART APEXII CCD area-detector
diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
 φ and ω scans
Absorption correction: multi-scan
(*SADABS*; Bruker, 2005)
 $T_{\min} = 0.958$, $T_{\max} = 0.980$

57259 measured reflections
13169 independent reflections
9552 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.032$
 $\theta_{\max} = 32.5^\circ$, $\theta_{\min} = 1.9^\circ$
 $h = -14 \rightarrow 14$
 $k = -18 \rightarrow 18$
 $l = -24 \rightarrow 24$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.058$
 $wR(F^2) = 0.169$
 $S = 1.02$
13169 reflections
528 parameters
144 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
 $w = 1/[\sigma^2(F_o^2) + (0.0817P)^2 + 0.7633P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.92 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.28 \text{ e } \text{\AA}^{-3}$

Special details

Experimental. The data was collected with the Oxford Cyrosystem Cobra low-temperature attachment.

Geometry. All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds 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 $F^2 > 2\text{sigma}(F^2)$ 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>	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
O1A	0.40902 (11)	0.86874 (9)	0.67186 (6)	0.0307 (2)	
O2A	-0.31593 (10)	0.42017 (8)	1.15549 (6)	0.02388 (18)	
O3A	0.30572 (13)	1.01757 (9)	0.60083 (7)	0.0387 (3)	
O4A	0.26641 (11)	0.94799 (9)	0.83284 (6)	0.0292 (2)	
C1A	-0.01076 (13)	0.73536 (10)	0.78085 (8)	0.0204 (2)	
H1AA	-0.0901	0.7821	0.7634	0.024*	
H1AB	-0.0321	0.6593	0.7789	0.024*	
C2A	0.12583 (13)	0.77666 (10)	0.71543 (8)	0.0213 (2)	
H2AA	0.2028	0.7253	0.7323	0.026*	
C3A	0.16635 (14)	0.89112 (11)	0.72390 (8)	0.0230 (2)	
H3AA	0.0878	0.9424	0.7101	0.028*	
C4A	0.18266 (13)	0.88549 (10)	0.81804 (8)	0.0222 (2)	
C5A	0.09136 (13)	0.80804 (10)	0.88799 (8)	0.0206 (2)	
H5AA	0.0960	0.8071	0.9471	0.025*	
C6A	-0.00023 (12)	0.73693 (10)	0.87393 (7)	0.0182 (2)	

C7A	-0.08885 (12)	0.65785 (10)	0.94693 (7)	0.0180 (2)
C8A	-0.05882 (13)	0.63354 (10)	1.03249 (8)	0.0209 (2)
H8AA	0.0167	0.6718	1.0439	0.025*
C9A	-0.13674 (13)	0.55552 (11)	1.09986 (8)	0.0220 (2)
H9AA	-0.1148	0.5408	1.1570	0.026*
C10A	-0.24827 (12)	0.49764 (10)	1.08465 (8)	0.0190 (2)
C11A	-0.28153 (13)	0.52069 (11)	1.00134 (8)	0.0214 (2)
H11A	-0.3580	0.4828	0.9906	0.026*
C12A	-0.20178 (13)	0.59991 (11)	0.93360 (8)	0.0215 (2)
H12A	-0.2248	0.6150	0.8767	0.026*
C13A	0.11848 (14)	0.77910 (11)	0.62083 (8)	0.0224 (2)
C14A	0.02075 (16)	0.84157 (12)	0.57676 (9)	0.0283 (3)
H14A	-0.0460	0.8836	0.6066	0.034*
C15A	0.01971 (17)	0.84323 (12)	0.48886 (9)	0.0314 (3)
H15A	-0.0477	0.8859	0.4594	0.038*
C16A	0.11732 (17)	0.78234 (13)	0.44497 (9)	0.0310 (3)
H16A	0.1164	0.7824	0.3856	0.037*
C17A	0.21553 (17)	0.72178 (14)	0.48779 (10)	0.0333 (3)
H17A	0.2833	0.6808	0.4575	0.040*
C18A	0.21615 (15)	0.72015 (12)	0.57507 (9)	0.0291 (3)
H18A	0.2846	0.6779	0.6038	0.035*
C19A	0.30060 (15)	0.93526 (11)	0.65901 (8)	0.0261 (3)
C20A	0.53784 (18)	0.89483 (17)	0.60642 (11)	0.0431 (4)
H20A	0.5832	0.9628	0.6106	0.052*
H20B	0.5168	0.9062	0.5466	0.052*
C21A	0.6333 (2)	0.7997 (2)	0.62443 (14)	0.0585 (6)
H21A	0.7196	0.8117	0.5789	0.088*
H21B	0.5845	0.7322	0.6241	0.088*
H21C	0.6585	0.7925	0.6821	0.088*
C22A	-0.42143 (15)	0.35266 (12)	1.14049 (9)	0.0284 (3)
H22A	-0.4546	0.2953	1.1941	0.043*
H22B	-0.3806	0.3184	1.0925	0.043*
H22C	-0.5015	0.3980	1.1245	0.043*
O1B	0.82102 (11)	0.36693 (9)	0.71886 (8)	0.0337 (2)
O2B	0.03578 (12)	-0.11336 (9)	1.25183 (7)	0.0331 (2)
O3B	0.67843 (13)	0.47429 (10)	0.79017 (8)	0.0387 (3)
O4B	0.77153 (12)	0.27422 (10)	0.92874 (8)	0.0373 (3)
C1B	0.38209 (13)	0.19261 (11)	0.88538 (8)	0.0232 (2)
H1BA	0.2799	0.2099	0.8943	0.028*
H1BB	0.3953	0.1312	0.8558	0.028*
C2B	0.46513 (14)	0.29275 (12)	0.82477 (9)	0.0260 (3)
H2BA	0.4347	0.3577	0.8493	0.031*
C3B	0.62513 (14)	0.27907 (11)	0.82326 (10)	0.0268 (3)
H3BA	0.6604	0.2205	0.7921	0.032*
C4B	0.65702 (15)	0.24696 (12)	0.91615 (10)	0.0283 (3)
C5B	0.55338 (14)	0.18059 (11)	0.98555 (9)	0.0256 (3)
H5BA	0.5774	0.1538	1.0418	0.031*
C6B	0.42349 (13)	0.15470 (10)	0.97426 (8)	0.0219 (2)

C7B	0.32038 (14)	0.08596 (10)	1.04606 (8)	0.0231 (2)
C8B	0.32756 (16)	0.07361 (12)	1.13529 (9)	0.0281 (3)
H8BA	0.3999	0.1117	1.1500	0.034*
C9B	0.23180 (17)	0.00737 (12)	1.20146 (9)	0.0306 (3)
H9BA	0.2384	0.0006	1.2611	0.037*
C10B	0.12475 (15)	-0.05007 (11)	1.18154 (9)	0.0263 (3)
C11B	0.11508 (14)	-0.03927 (12)	1.09398 (9)	0.0262 (3)
H11B	0.0428	-0.0778	1.0796	0.031*
C12B	0.21182 (14)	0.02814 (11)	1.02784 (9)	0.0248 (2)
H12B	0.2042	0.0353	0.9683	0.030*
C13B	0.4318 (8)	0.3146 (4)	0.7363 (6)	0.0160 (10) 0.545 (17)
C14B	0.3759 (8)	0.4132 (4)	0.6948 (3)	0.0288 (9) 0.545 (17)
H14B	0.3605	0.4707	0.7246	0.035* 0.545 (17)
C15B	0.3424 (9)	0.4304 (4)	0.6127 (3)	0.0367 (11) 0.545 (17)
H15B	0.3050	0.4992	0.5877	0.044* 0.545 (17)
C16B	0.3613 (9)	0.3510 (6)	0.5657 (5)	0.0338 (12) 0.545 (17)
H16B	0.3358	0.3663	0.5094	0.041* 0.545 (17)
C17B	0.4160 (12)	0.2501 (7)	0.5978 (5)	0.0249 (11) 0.545 (17)
H17B	0.4367	0.1944	0.5664	0.030* 0.545 (17)
C18B	0.4363 (14)	0.2418 (9)	0.6820 (6)	0.035 (2) 0.545 (17)
H18B	0.4587	0.1693	0.7101	0.042* 0.545 (17)
C13C	0.4419 (11)	0.3148 (7)	0.7251 (8)	0.033 (2) 0.455 (17)
C14C	0.4258 (11)	0.4248 (6)	0.6876 (4)	0.0334 (13) 0.455 (17)
H14C	0.4284	0.4795	0.7194	0.040* 0.455 (17)
C15C	0.4053 (12)	0.4520 (6)	0.6006 (4)	0.0403 (16) 0.455 (17)
H15C	0.3968	0.5268	0.5711	0.048* 0.455 (17)
C16C	0.3975 (10)	0.3702 (8)	0.5579 (6)	0.0361 (16) 0.455 (17)
H16C	0.3833	0.3855	0.4989	0.043* 0.455 (17)
C17C	0.4117 (16)	0.2646 (9)	0.6059 (7)	0.037 (2) 0.455 (17)
H17C	0.3922	0.2086	0.5788	0.044* 0.455 (17)
C18C	0.4503 (14)	0.2261 (9)	0.6885 (6)	0.0201 (12) 0.455 (17)
H18C	0.4767	0.1535	0.7137	0.024* 0.455 (17)
C19B	0.70784 (15)	0.38573 (12)	0.77640 (10)	0.0288 (3)
C20B	0.91687 (18)	0.45985 (15)	0.67563 (13)	0.0416 (4)
H20C	1.0112	0.4325	0.6538	0.050*
H20D	0.9281	0.5022	0.7189	0.050*
C21B	0.8642 (2)	0.53375 (16)	0.60068 (12)	0.0456 (4)
H21D	0.9318	0.5951	0.5732	0.068*
H21E	0.7718	0.5623	0.6223	0.068*
H21F	0.8543	0.4924	0.5573	0.068*
C22B	-0.06909 (17)	-0.18023 (14)	1.23449 (11)	0.0366 (3)
H22D	-0.1213	-0.2252	1.2902	0.055*
H22E	-0.0223	-0.2283	1.1980	0.055*
H22F	-0.1352	-0.1328	1.2034	0.055*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1A	0.0293 (5)	0.0386 (6)	0.0216 (4)	-0.0049 (4)	-0.0003 (4)	-0.0053 (4)
O2A	0.0278 (4)	0.0217 (4)	0.0209 (4)	-0.0044 (3)	-0.0052 (3)	-0.0017 (3)
O3A	0.0524 (7)	0.0269 (5)	0.0314 (6)	-0.0069 (5)	-0.0031 (5)	0.0001 (4)
O4A	0.0363 (5)	0.0302 (5)	0.0228 (4)	-0.0127 (4)	-0.0039 (4)	-0.0094 (4)
C1A	0.0222 (5)	0.0234 (6)	0.0168 (5)	-0.0032 (4)	-0.0050 (4)	-0.0059 (4)
C2A	0.0248 (5)	0.0220 (5)	0.0178 (5)	-0.0019 (4)	-0.0046 (4)	-0.0057 (4)
C3A	0.0283 (6)	0.0218 (6)	0.0195 (5)	-0.0026 (5)	-0.0042 (4)	-0.0062 (4)
C4A	0.0262 (6)	0.0233 (6)	0.0183 (5)	-0.0033 (5)	-0.0033 (4)	-0.0079 (4)
C5A	0.0233 (5)	0.0238 (6)	0.0156 (5)	-0.0029 (4)	-0.0027 (4)	-0.0067 (4)
C6A	0.0182 (5)	0.0204 (5)	0.0171 (5)	0.0012 (4)	-0.0042 (4)	-0.0061 (4)
C7A	0.0171 (5)	0.0206 (5)	0.0170 (5)	0.0009 (4)	-0.0037 (4)	-0.0056 (4)
C8A	0.0207 (5)	0.0239 (6)	0.0200 (5)	-0.0018 (4)	-0.0066 (4)	-0.0067 (4)
C9A	0.0244 (5)	0.0246 (6)	0.0181 (5)	-0.0004 (4)	-0.0078 (4)	-0.0045 (4)
C10A	0.0197 (5)	0.0179 (5)	0.0192 (5)	0.0014 (4)	-0.0038 (4)	-0.0041 (4)
C11A	0.0193 (5)	0.0248 (6)	0.0210 (5)	-0.0024 (4)	-0.0061 (4)	-0.0049 (4)
C12A	0.0198 (5)	0.0263 (6)	0.0191 (5)	-0.0016 (4)	-0.0061 (4)	-0.0046 (4)
C13A	0.0278 (6)	0.0233 (6)	0.0168 (5)	-0.0059 (5)	-0.0049 (4)	-0.0051 (4)
C14A	0.0340 (7)	0.0278 (6)	0.0248 (6)	0.0019 (5)	-0.0084 (5)	-0.0078 (5)
C15A	0.0421 (8)	0.0286 (7)	0.0237 (6)	-0.0020 (6)	-0.0129 (6)	-0.0015 (5)
C16A	0.0419 (8)	0.0354 (7)	0.0163 (5)	-0.0095 (6)	-0.0061 (5)	-0.0055 (5)
C17A	0.0356 (7)	0.0424 (8)	0.0255 (6)	-0.0003 (6)	-0.0043 (5)	-0.0164 (6)
C18A	0.0326 (7)	0.0345 (7)	0.0243 (6)	0.0012 (6)	-0.0094 (5)	-0.0118 (5)
C19A	0.0341 (7)	0.0251 (6)	0.0190 (5)	-0.0091 (5)	-0.0016 (5)	-0.0067 (5)
C20A	0.0326 (7)	0.0604 (11)	0.0305 (8)	-0.0117 (7)	0.0078 (6)	-0.0094 (7)
C21A	0.0303 (8)	0.0938 (17)	0.0459 (11)	0.0043 (9)	0.0024 (7)	-0.0146 (11)
C22A	0.0315 (6)	0.0246 (6)	0.0281 (6)	-0.0078 (5)	-0.0072 (5)	-0.0025 (5)
O1B	0.0325 (5)	0.0284 (5)	0.0403 (6)	-0.0011 (4)	-0.0040 (4)	-0.0110 (4)
O2B	0.0379 (5)	0.0327 (5)	0.0260 (5)	0.0015 (4)	-0.0083 (4)	-0.0003 (4)
O3B	0.0427 (6)	0.0348 (6)	0.0435 (6)	0.0031 (5)	-0.0090 (5)	-0.0181 (5)
O4B	0.0319 (5)	0.0417 (6)	0.0440 (6)	-0.0052 (5)	-0.0215 (5)	-0.0093 (5)
C1B	0.0226 (5)	0.0267 (6)	0.0234 (6)	0.0011 (5)	-0.0094 (4)	-0.0083 (5)
C2B	0.0253 (6)	0.0276 (6)	0.0289 (6)	0.0023 (5)	-0.0093 (5)	-0.0109 (5)
C3B	0.0252 (6)	0.0259 (6)	0.0338 (7)	0.0018 (5)	-0.0103 (5)	-0.0124 (5)
C4B	0.0260 (6)	0.0295 (7)	0.0344 (7)	0.0026 (5)	-0.0146 (5)	-0.0108 (5)
C5B	0.0294 (6)	0.0249 (6)	0.0280 (6)	0.0044 (5)	-0.0144 (5)	-0.0105 (5)
C6B	0.0257 (6)	0.0210 (5)	0.0234 (6)	0.0051 (4)	-0.0096 (4)	-0.0101 (4)
C7B	0.0279 (6)	0.0220 (6)	0.0232 (6)	0.0055 (5)	-0.0097 (5)	-0.0093 (5)
C8B	0.0376 (7)	0.0260 (6)	0.0254 (6)	0.0020 (5)	-0.0142 (5)	-0.0088 (5)
C9B	0.0426 (8)	0.0284 (7)	0.0236 (6)	0.0045 (6)	-0.0137 (6)	-0.0067 (5)
C10B	0.0305 (6)	0.0247 (6)	0.0237 (6)	0.0080 (5)	-0.0079 (5)	-0.0046 (5)
C11B	0.0258 (6)	0.0289 (6)	0.0263 (6)	0.0039 (5)	-0.0083 (5)	-0.0088 (5)
C12B	0.0259 (6)	0.0285 (6)	0.0230 (6)	0.0041 (5)	-0.0089 (5)	-0.0093 (5)
C13B	0.0167 (15)	0.0147 (16)	0.017 (2)	0.0025 (12)	-0.0014 (12)	-0.0057 (12)
C14B	0.031 (2)	0.0253 (15)	0.0305 (15)	-0.0005 (16)	-0.0052 (16)	-0.0084 (11)
C15B	0.049 (3)	0.0263 (16)	0.0327 (17)	0.0033 (17)	-0.0118 (18)	-0.0008 (12)

C16B	0.043 (3)	0.036 (2)	0.0238 (19)	-0.005 (2)	-0.013 (2)	-0.0042 (15)
C17B	0.042 (2)	0.0218 (18)	0.0146 (16)	-0.0060 (16)	-0.0051 (15)	-0.0108 (18)
C18B	0.049 (4)	0.018 (3)	0.035 (3)	0.001 (2)	-0.014 (2)	0.0042 (18)
C13C	0.030 (3)	0.055 (4)	0.015 (3)	-0.017 (2)	-0.008 (2)	-0.003 (2)
C14C	0.040 (3)	0.035 (2)	0.028 (2)	0.006 (2)	-0.011 (2)	-0.0114 (17)
C15C	0.054 (4)	0.035 (2)	0.031 (2)	0.014 (3)	-0.012 (2)	-0.0056 (17)
C16C	0.033 (3)	0.050 (4)	0.026 (2)	0.005 (2)	-0.012 (2)	-0.006 (3)
C17C	0.045 (3)	0.041 (4)	0.032 (3)	-0.016 (3)	0.002 (2)	-0.028 (2)
C18C	0.033 (2)	0.012 (3)	0.017 (2)	0.0041 (19)	-0.0100 (16)	-0.0024 (19)
C19B	0.0255 (6)	0.0328 (7)	0.0323 (7)	-0.0010 (5)	-0.0109 (5)	-0.0116 (6)
C20B	0.0308 (7)	0.0402 (9)	0.0492 (10)	-0.0045 (6)	-0.0038 (7)	-0.0047 (7)
C21B	0.0490 (10)	0.0414 (9)	0.0433 (9)	-0.0057 (8)	-0.0069 (8)	-0.0050 (7)
C22B	0.0342 (7)	0.0356 (8)	0.0353 (8)	-0.0006 (6)	-0.0090 (6)	0.0021 (6)

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

O1A—C19A	1.3320 (18)	C1B—C6B	1.5092 (17)
O1A—C20A	1.4452 (18)	C1B—C2B	1.5220 (19)
O2A—C10A	1.3648 (15)	C1B—H1BA	0.99
O2A—C22A	1.4289 (16)	C1B—H1BB	0.99
O3A—C19A	1.2038 (17)	C2B—C13B	1.468 (8)
O4A—C4A	1.2313 (15)	C2B—C3B	1.5385 (19)
C1A—C6A	1.5143 (16)	C2B—C13C	1.607 (12)
C1A—C2A	1.5257 (17)	C2B—H2BA	1.00
C1A—H1AA	0.99	C3B—C19B	1.524 (2)
C1A—H1AB	0.99	C3B—C4B	1.528 (2)
C2A—C13A	1.5189 (16)	C3B—H3BA	1.00
C2A—C3A	1.5412 (17)	C4B—C5B	1.444 (2)
C2A—H2AA	1.00	C5B—C6B	1.3541 (18)
C3A—C19A	1.5131 (18)	C5B—H5BA	0.95
C3A—C4A	1.5259 (17)	C6B—C7B	1.4723 (19)
C3A—H3AA	1.00	C7B—C12B	1.4000 (18)
C4A—C5A	1.4498 (17)	C7B—C8B	1.4104 (18)
C5A—C6A	1.3554 (16)	C8B—C9B	1.376 (2)
C5A—H5AA	0.95	C8B—H8BA	0.95
C6A—C7A	1.4726 (16)	C9B—C10B	1.399 (2)
C7A—C12A	1.3997 (16)	C9B—H9BA	0.95
C7A—C8A	1.4099 (16)	C10B—C11B	1.3927 (18)
C8A—C9A	1.3757 (18)	C11B—C12B	1.387 (2)
C8A—H8AA	0.95	C11B—H11B	0.95
C9A—C10A	1.4013 (17)	C12B—H12B	0.95
C9A—H9AA	0.95	C13B—C18B	1.398 (8)
C10A—C11A	1.3884 (16)	C13B—C14B	1.399 (6)
C11A—C12A	1.3941 (17)	C14B—C15B	1.375 (6)
C11A—H11A	0.95	C14B—H14B	0.95
C12A—H12A	0.95	C15B—C16B	1.373 (7)
C13A—C18A	1.3861 (19)	C15B—H15B	0.95
C13A—C14A	1.3917 (19)	C16B—C17B	1.376 (8)

C14A—C15A	1.4008 (19)	C16B—H16B	0.95
C14A—H14A	0.95	C17B—C18B	1.376 (8)
C15A—C16A	1.386 (2)	C17B—H17B	0.95
C15A—H15A	0.95	C18B—H18B	0.95
C16A—C17A	1.375 (2)	C13C—C18C	1.371 (9)
C16A—H16A	0.95	C13C—C14C	1.381 (9)
C17A—C18A	1.3903 (19)	C14C—C15C	1.399 (7)
C17A—H17A	0.95	C14C—H14C	0.95
C18A—H18A	0.95	C15C—C16C	1.374 (8)
C20A—C21A	1.491 (3)	C15C—H15C	0.95
C20A—H20A	0.99	C16C—C17C	1.377 (10)
C20A—H20B	0.99	C16C—H16C	0.95
C21A—H21A	0.98	C17C—C18C	1.407 (10)
C21A—H21B	0.98	C17C—H17C	0.95
C21A—H21C	0.98	C18C—H18C	0.95
C22A—H22A	0.98	C20B—C21B	1.488 (3)
C22A—H22B	0.98	C20B—H20C	0.99
C22A—H22C	0.98	C20B—H20D	0.99
O1B—C19B	1.3373 (18)	C21B—H21D	0.98
O1B—C20B	1.451 (2)	C21B—H21E	0.98
O2B—C10B	1.3619 (18)	C21B—H21F	0.98
O2B—C22B	1.436 (2)	C22B—H22D	0.98
O3B—C19B	1.1995 (18)	C22B—H22E	0.98
O4B—C4B	1.2279 (16)	C22B—H22F	0.98
C19A—O1A—C20A	116.22 (12)	C13B—C2B—C13C	4.7 (6)
C10A—O2A—C22A	117.00 (10)	C1B—C2B—C13C	113.0 (4)
C6A—C1A—C2A	112.11 (9)	C3B—C2B—C13C	107.6 (4)
C6A—C1A—H1AA	109.2	C13B—C2B—H2BA	107.6
C2A—C1A—H1AA	109.2	C1B—C2B—H2BA	107.6
C6A—C1A—H1AB	109.2	C3B—C2B—H2BA	107.6
C2A—C1A—H1AB	109.2	C13C—C2B—H2BA	110.3
H1AA—C1A—H1AB	107.9	C19B—C3B—C4B	106.55 (11)
C13A—C2A—C1A	113.81 (10)	C19B—C3B—C2B	111.56 (11)
C13A—C2A—C3A	110.83 (10)	C4B—C3B—C2B	111.59 (12)
C1A—C2A—C3A	109.34 (10)	C19B—C3B—H3BA	109.0
C13A—C2A—H2AA	107.5	C4B—C3B—H3BA	109.0
C1A—C2A—H2AA	107.5	C2B—C3B—H3BA	109.0
C3A—C2A—H2AA	107.5	O4B—C4B—C5B	122.18 (13)
C19A—C3A—C4A	111.12 (10)	O4B—C4B—C3B	119.32 (14)
C19A—C3A—C2A	110.74 (10)	C5B—C4B—C3B	118.42 (11)
C4A—C3A—C2A	110.11 (10)	C6B—C5B—C4B	123.11 (12)
C19A—C3A—H3AA	108.3	C6B—C5B—H5BA	118.4
C4A—C3A—H3AA	108.3	C4B—C5B—H5BA	118.4
C2A—C3A—H3AA	108.3	C5B—C6B—C7B	121.91 (11)
O4A—C4A—C5A	122.27 (11)	C5B—C6B—C1B	120.34 (12)
O4A—C4A—C3A	120.41 (11)	C7B—C6B—C1B	117.70 (11)
C5A—C4A—C3A	117.25 (10)	C12B—C7B—C8B	117.19 (12)

C6A—C5A—C4A	123.75 (11)	C12B—C7B—C6B	120.73 (11)
C6A—C5A—H5AA	118.1	C8B—C7B—C6B	122.07 (12)
C4A—C5A—H5AA	118.1	C9B—C8B—C7B	121.25 (13)
C5A—C6A—C7A	121.98 (10)	C9B—C8B—H8BA	119.4
C5A—C6A—C1A	119.92 (11)	C7B—C8B—H8BA	119.4
C7A—C6A—C1A	118.08 (10)	C8B—C9B—C10B	120.48 (12)
C12A—C7A—C8A	117.23 (11)	C8B—C9B—H9BA	119.8
C12A—C7A—C6A	121.58 (10)	C10B—C9B—H9BA	119.8
C8A—C7A—C6A	121.12 (10)	O2B—C10B—C11B	124.76 (13)
C9A—C8A—C7A	121.32 (11)	O2B—C10B—C9B	115.76 (12)
C9A—C8A—H8AA	119.3	C11B—C10B—C9B	119.48 (13)
C7A—C8A—H8AA	119.3	C12B—C11B—C10B	119.52 (12)
C8A—C9A—C10A	120.40 (11)	C12B—C11B—H11B	120.2
C8A—C9A—H9AA	119.8	C10B—C11B—H11B	120.2
C10A—C9A—H9AA	119.8	C11B—C12B—C7B	122.08 (12)
O2A—C10A—C11A	124.88 (11)	C11B—C12B—H12B	119.0
O2A—C10A—C9A	115.49 (10)	C7B—C12B—H12B	119.0
C11A—C10A—C9A	119.62 (11)	C18B—C13B—C14B	107.6 (8)
C10A—C11A—C12A	119.44 (11)	C18B—C13B—C2B	128.1 (5)
C10A—C11A—H11A	120.3	C14B—C13B—C2B	124.1 (5)
C12A—C11A—H11A	120.3	C15B—C14B—C13B	122.8 (5)
C11A—C12A—C7A	121.98 (11)	C15B—C14B—H14B	118.6
C11A—C12A—H12A	119.0	C13B—C14B—H14B	118.6
C7A—C12A—H12A	119.0	C16B—C15B—C14B	122.0 (5)
C18A—C13A—C14A	118.29 (12)	C16B—C15B—H15B	119.0
C18A—C13A—C2A	118.99 (12)	C14B—C15B—H15B	119.0
C14A—C13A—C2A	122.68 (12)	C15B—C16B—C17B	121.9 (7)
C13A—C14A—C15A	120.77 (13)	C15B—C16B—H16B	119.1
C13A—C14A—H14A	119.6	C17B—C16B—H16B	119.1
C15A—C14A—H14A	119.6	C18B—C17B—C16B	109.9 (9)
C16A—C15A—C14A	119.77 (13)	C18B—C17B—H17B	125.1
C16A—C15A—H15A	120.1	C16B—C17B—H17B	125.1
C14A—C15A—H15A	120.1	C17B—C18B—C13B	135.3 (10)
C17A—C16A—C15A	119.70 (12)	C17B—C18B—H18B	112.4
C17A—C16A—H16A	120.1	C13B—C18B—H18B	112.4
C15A—C16A—H16A	120.1	C18C—C13C—C14C	130.0 (11)
C16A—C17A—C18A	120.39 (14)	C18C—C13C—C2B	117.1 (7)
C16A—C17A—H17A	119.8	C14C—C13C—C2B	112.7 (7)
C18A—C17A—H17A	119.8	C13C—C14C—C15C	116.8 (7)
C13A—C18A—C17A	121.06 (13)	C13C—C14C—H14C	121.6
C13A—C18A—H18A	119.5	C15C—C14C—H14C	121.6
C17A—C18A—H18A	119.5	C16C—C15C—C14C	119.8 (6)
O3A—C19A—O1A	124.84 (13)	C16C—C15C—H15C	120.1
O3A—C19A—C3A	123.63 (14)	C14C—C15C—H15C	120.1
O1A—C19A—C3A	111.44 (11)	C15C—C16C—C17C	115.8 (8)
O1A—C20A—C21A	106.78 (15)	C15C—C16C—H16C	122.1
O1A—C20A—H20A	110.4	C17C—C16C—H16C	122.1
C21A—C20A—H20A	110.4	C16C—C17C—C18C	130.6 (11)

O1A—C20A—H20B	110.4	C16C—C17C—H17C	114.7
C21A—C20A—H20B	110.4	C18C—C17C—H17C	114.7
H20A—C20A—H20B	108.6	C13C—C18C—C17C	105.8 (11)
C20A—C21A—H21A	109.5	C13C—C18C—H18C	127.1
C20A—C21A—H21B	109.5	C17C—C18C—H18C	127.1
H21A—C21A—H21B	109.5	O3B—C19B—O1B	124.16 (14)
C20A—C21A—H21C	109.5	O3B—C19B—C3B	125.19 (14)
H21A—C21A—H21C	109.5	O1B—C19B—C3B	110.63 (12)
H21B—C21A—H21C	109.5	O1B—C20B—C21B	111.94 (14)
O2A—C22A—H22A	109.5	O1B—C20B—H20C	109.2
O2A—C22A—H22B	109.5	C21B—C20B—H20C	109.2
H22A—C22A—H22B	109.5	O1B—C20B—H20D	109.2
O2A—C22A—H22C	109.5	C21B—C20B—H20D	109.2
H22A—C22A—H22C	109.5	H20C—C20B—H20D	107.9
H22B—C22A—H22C	109.5	C20B—C21B—H21D	109.5
C19B—O1B—C20B	116.36 (12)	C20B—C21B—H21E	109.5
C10B—O2B—C22B	117.72 (11)	H21D—C21B—H21E	109.5
C6B—C1B—C2B	114.71 (10)	C20B—C21B—H21F	109.5
C6B—C1B—H1BA	108.6	H21D—C21B—H21F	109.5
C2B—C1B—H1BA	108.6	H21E—C21B—H21F	109.5
C6B—C1B—H1BB	108.6	O2B—C22B—H22D	109.5
C2B—C1B—H1BB	108.6	O2B—C22B—H22E	109.5
H1BA—C1B—H1BB	107.6	H22D—C22B—H22E	109.5
C13B—C2B—C1B	110.9 (3)	O2B—C22B—H22F	109.5
C13B—C2B—C3B	112.2 (3)	H22D—C22B—H22F	109.5
C1B—C2B—C3B	110.68 (11)	H22E—C22B—H22F	109.5
C6A—C1A—C2A—C13A	-178.56 (10)	C2B—C3B—C4B—O4B	-151.15 (14)
C6A—C1A—C2A—C3A	-54.04 (13)	C19B—C3B—C4B—C5B	153.85 (12)
C13A—C2A—C3A—C19A	-53.03 (14)	C2B—C3B—C4B—C5B	31.85 (17)
C1A—C2A—C3A—C19A	-179.27 (10)	O4B—C4B—C5B—C6B	176.25 (14)
C13A—C2A—C3A—C4A	-176.33 (10)	C3B—C4B—C5B—C6B	-6.8 (2)
C1A—C2A—C3A—C4A	57.43 (13)	C4B—C5B—C6B—C7B	179.42 (12)
C19A—C3A—C4A—O4A	26.66 (17)	C4B—C5B—C6B—C1B	1.78 (19)
C2A—C3A—C4A—O4A	149.73 (12)	C2B—C1B—C6B—C5B	-22.94 (17)
C19A—C3A—C4A—C5A	-156.21 (12)	C2B—C1B—C6B—C7B	159.32 (11)
C2A—C3A—C4A—C5A	-33.14 (15)	C5B—C6B—C7B—C12B	-157.01 (12)
O4A—C4A—C5A—C6A	-178.87 (13)	C1B—C6B—C7B—C12B	20.69 (17)
C3A—C4A—C5A—C6A	4.06 (19)	C5B—C6B—C7B—C8B	21.79 (19)
C4A—C5A—C6A—C7A	178.62 (11)	C1B—C6B—C7B—C8B	-160.50 (12)
C4A—C5A—C6A—C1A	0.20 (18)	C12B—C7B—C8B—C9B	0.1 (2)
C2A—C1A—C6A—C5A	25.66 (16)	C6B—C7B—C8B—C9B	-178.77 (13)
C2A—C1A—C6A—C7A	-152.81 (10)	C7B—C8B—C9B—C10B	0.3 (2)
C5A—C6A—C7A—C12A	167.45 (12)	C22B—O2B—C10B—C11B	4.9 (2)
C1A—C6A—C7A—C12A	-14.11 (16)	C22B—O2B—C10B—C9B	-175.43 (12)
C5A—C6A—C7A—C8A	-15.68 (18)	C8B—C9B—C10B—O2B	179.92 (12)
C1A—C6A—C7A—C8A	162.76 (11)	C8B—C9B—C10B—C11B	-0.4 (2)
C12A—C7A—C8A—C9A	0.45 (18)	O2B—C10B—C11B—C12B	179.76 (12)

C6A—C7A—C8A—C9A	−176.55 (11)	C9B—C10B—C11B—C12B	0.1 (2)
C7A—C8A—C9A—C10A	0.29 (19)	C10B—C11B—C12B—C7B	0.3 (2)
C22A—O2A—C10A—C11A	4.94 (17)	C8B—C7B—C12B—C11B	−0.37 (19)
C22A—O2A—C10A—C9A	−174.41 (11)	C6B—C7B—C12B—C11B	178.49 (12)
C8A—C9A—C10A—O2A	178.32 (11)	C1B—C2B—C13B—C18B	−52.5 (10)
C8A—C9A—C10A—C11A	−1.07 (18)	C3B—C2B—C13B—C18B	71.8 (9)
O2A—C10A—C11A—C12A	−178.25 (11)	C13C—C2B—C13B—C18B	65 (6)
C9A—C10A—C11A—C12A	1.08 (18)	C1B—C2B—C13B—C14B	121.2 (6)
C10A—C11A—C12A—C7A	−0.33 (19)	C3B—C2B—C13B—C14B	−114.5 (6)
C8A—C7A—C12A—C11A	−0.43 (18)	C13C—C2B—C13B—C14B	−122 (7)
C6A—C7A—C12A—C11A	176.55 (11)	C18B—C13B—C14B—C15B	−3.5 (9)
C1A—C2A—C13A—C18A	−122.41 (13)	C2B—C13B—C14B—C15B	−178.3 (5)
C3A—C2A—C13A—C18A	113.87 (14)	C13B—C14B—C15B—C16B	0.2 (7)
C1A—C2A—C13A—C14A	60.06 (16)	C14B—C15B—C16B—C17B	−0.6 (9)
C3A—C2A—C13A—C14A	−63.66 (16)	C15B—C16B—C17B—C18B	4.2 (13)
C18A—C13A—C14A—C15A	1.1 (2)	C16B—C17B—C18B—C13B	−10 (2)
C2A—C13A—C14A—C15A	178.63 (12)	C14B—C13B—C18B—C17B	9.5 (19)
C13A—C14A—C15A—C16A	−0.2 (2)	C2B—C13B—C18B—C17B	−176.0 (12)
C14A—C15A—C16A—C17A	−0.8 (2)	C13B—C2B—C13C—C18C	−111 (7)
C15A—C16A—C17A—C18A	0.9 (2)	C1B—C2B—C13C—C18C	−46.3 (10)
C14A—C13A—C18A—C17A	−1.0 (2)	C3B—C2B—C13C—C18C	76.2 (9)
C2A—C13A—C18A—C17A	−178.61 (13)	C13B—C2B—C13C—C14C	73 (6)
C16A—C17A—C18A—C13A	0.0 (2)	C1B—C2B—C13C—C14C	137.9 (6)
C20A—O1A—C19A—O3A	−4.8 (2)	C3B—C2B—C13C—C14C	−99.7 (7)
C20A—O1A—C19A—C3A	171.95 (12)	C18C—C13C—C14C—C15C	4.7 (13)
C4A—C3A—C19A—O3A	−119.90 (14)	C2B—C13C—C14C—C15C	179.9 (5)
C2A—C3A—C19A—O3A	117.39 (14)	C13C—C14C—C15C—C16C	2.1 (8)
C4A—C3A—C19A—O1A	63.32 (14)	C14C—C15C—C16C—C17C	−0.3 (11)
C2A—C3A—C19A—O1A	−59.39 (14)	C15C—C16C—C17C—C18C	−8.9 (19)
C19A—O1A—C20A—C21A	−169.57 (14)	C14C—C13C—C18C—C17C	−11.1 (17)
C6B—C1B—C2B—C13B	172.3 (3)	C2B—C13C—C18C—C17C	173.9 (9)
C6B—C1B—C2B—C3B	47.14 (15)	C16C—C17C—C18C—C13C	13 (2)
C6B—C1B—C2B—C13C	167.8 (4)	C20B—O1B—C19B—O3B	4.1 (2)
C13B—C2B—C3B—C19B	65.7 (2)	C20B—O1B—C19B—C3B	−174.31 (12)
C1B—C2B—C3B—C19B	−169.88 (11)	C4B—C3B—C19B—O3B	−73.10 (17)
C13C—C2B—C3B—C19B	66.3 (3)	C2B—C3B—C19B—O3B	48.92 (19)
C13B—C2B—C3B—C4B	−175.3 (2)	C4B—C3B—C19B—O1B	105.25 (13)
C1B—C2B—C3B—C4B	−50.81 (14)	C2B—C3B—C19B—O1B	−132.73 (12)
C13C—C2B—C3B—C4B	−174.7 (3)	C19B—O1B—C20B—C21B	−80.67 (19)
C19B—C3B—C4B—O4B	−29.15 (18)		

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

$D\text{—H}\cdots A$	$D\text{—H}$	$H\cdots A$	$D\cdots A$	$D\text{—H}\cdots A$
C3A—H3A _A —O2B ⁱ	1.00	2.57	3.4093 (18)	142
C5A—H5A _A —O4B ⁱⁱ	0.95	2.55	3.3630 (17)	143
C8A—H8A _A —O4B ⁱⁱ	0.95	2.31	3.2446 (18)	166
C14B—H14B—O2A ⁱ	0.95	2.58	3.500 (5)	163

C16A—H16A···O2B ⁱⁱⁱ	0.95	2.47	3.2789 (18)	143
C17B—H17B···O3A ^{iv}	0.95	2.47	3.116 (9)	125
C21B—H21D···Cg1 ^v	0.98	2.85	3.811 (2)	167
C22B—H22D···Cg2 ^{vi}	0.98	2.87	3.644 (4)	137

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