

6,8-Dichloro-4-oxochromene-3-carbaldehyde

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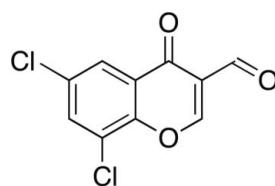
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Key indicators: single-crystal X-ray study; $T = 100\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.010\text{ \AA}$; R factor = 0.076; wR factor = 0.212; data-to-parameter ratio = 15.5.

The asymmetric unit of the title compound, $\text{C}_{10}\text{H}_4\text{Cl}_2\text{O}_3$, contain two essentially planar independent molecules (mean atomic deviations from the corresponding least-square planes are 0.041 and 0.045 Å for molecules 1 and 2, respectively). In the crystal, molecules are linked through a pair of halogen bonds [$\text{Cl} \cdots \text{O}$ separations are 3.044 (5) and 3.033 (6) Å, $\text{C}-\text{Cl} \cdots \text{O}$ angles are 160.4 (3) and 162.8 (3)°, and $\text{C}=\text{O} \cdots \text{Cl}$ angles are 138.7 (4) and 139.6 (4)°, respectively, in molecules 1 and 2] and $\text{C}-\text{H} \cdots \text{O}$ hydrogen bonds into slightly folded bands [the dihedral angle between the planes of neighboring molecules is 8.6 (2)°] along the c -axis direction.

Related literature

For the biological activity of the title and related compounds, see: Shim *et al.* (2003); Kawase *et al.* (2007); Dückert *et al.* (2012). For related structures, see: Ishikawa *et al.* (2013a,b). For halogen bonding, see: Auffinger *et al.* (2004); Metrangolo *et al.* (2005); Wilcken *et al.* (2013).



Experimental

Crystal data

$\text{C}_{10}\text{H}_4\text{Cl}_2\text{O}_3$	$\alpha = 96.55$ (6)°
$M_r = 243.05$	$\beta = 92.23$ (7)°
Triclinic, $P\bar{1}$	$\gamma = 101.98$ (7)°
$a = 8.288$ (8) Å	$V = 917.2$ (13) Å ³
$b = 8.325$ (7) Å	$Z = 4$
$c = 13.706$ (7) Å	Mo $K\alpha$ radiation

$\mu = 0.68\text{ mm}^{-1}$
 $T = 100\text{ K}$

$0.42 \times 0.22 \times 0.08\text{ mm}$

Data collection

Rigaku AFC-7R diffractometer
Absorption correction: ψ scan
(North *et al.*, 1968)
 $T_{\min} = 0.841$, $T_{\max} = 0.947$
5130 measured reflections
4203 independent reflections

2596 reflections with $F^2 > 2\sigma(F^2)$
 $R_{\text{int}} = 0.057$
3 standard reflections every 150
reflections
intensity decay: 4.9%

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.076$
 $wR(F^2) = 0.212$
 $S = 1.10$
4203 reflections

271 parameters
H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.70\text{ e \AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.79\text{ e \AA}^{-3}$

Table 1
Hydrogen-bond geometry (Å, °).

$D-\text{H} \cdots A$	$D-\text{H}$	$\text{H} \cdots A$	$D \cdots A$	$D-\text{H} \cdots A$
$\text{C}4b^i - \text{H}2b^i \cdots \text{O}2a$	0.95	2.35	3.246 (8)	157
$\text{C}4a - \text{H}2a \cdots \text{O}2b^i$	0.95	2.35	3.259 (8)	160

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

Data collection: *WinAFC* (Rigaku, 1999); cell refinement: *WinAFC*; data reduction: *WinAFC*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *CrystalStructure* (Rigaku, 2010); software used to prepare material for publication: *CrystalStructure*.

We acknowledge the University of Shizuoka for instrumental support.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: LD2111).

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supplementary materials

Acta Cryst. (2013). E69, o1416 [doi:10.1107/S1600536813022228]

6,8-Dichloro-4-oxochromene-3-carbaldehyde

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1. Comment

6,8-Dichloro-3-formylchromone shows many biological functions such as protein tyrosine phosphatase inhibitory (Shim *et al.* 2003), tumor cell-cytotoxic, anti-HIV, anti-*Helicobacter pylori*, and urease inhibitory activities (Kawase *et al.* 2007). In addition, it is used as a starting material for the synthesis of biologically relevant molecules (Dückert *et al.* 2012).

The title compound, $C_{10}H_4Cl_2O_3$, crystallizes with two independent molecules in the asymmetric unit (Fig. 1). The mean deviations from the least-square planes for all atoms of molecule 1 and 2 are 0.0410 Å and 0.0449 Å, respectively. In addition, the largest deviations of molecule 1 and 2 are 0.1512 (17) Å for Cl1a and -0.0973 Å for H4b, respectively. This means that all atoms of each molecule are essentially coplanar.

In the crystal, the molecules 1 and 2 are linked to each other through intermolecular interactions of the Cl atoms at the 8-position with the O atoms of the formyl groups [$Cl2a\cdots O3b^i$; 3.033 (6) Å, $Cl2b^i\cdots O3a$; 3.044 (5) Å, $C7a-Cl2a\cdots O3b^i$ = 160.4 (3)°, $C7b^i-Cl2b^i\cdots O3a$ = 162.8 (3)°, $C10a-O3a\cdots Cl2b^i$ = 138.7 (4)°, $C10b^i-O3b^i\cdots Cl2a$ = 139.6 (4)° (i): $-x + 1, -y + 1, -z + 1$], and the carbonyl O atoms at the 4-position with the C–H atoms at the 5-position. The short contacts and the geometries involved in the Cl atoms fall into halogen bonding (Auffinger *et al.* 2004). Due to these halogen and hydrogen bonds, the molecules form wavy bands along *c* axis, as shown in Fig. 2.

Halogen bonds have been found to occur in organic, inorganic, and biological systems, and have recently attracted much attention in medicinal chemistry, chemical biology, and supramolecular chemistry (Auffinger *et al.* 2004, Metrangolo *et al.* 2005, Wilcken *et al.* 2013). Our analysis suggests that the strong inhibitory activity of the title compound against urease may be attributable to the halogen bond observed in the crystal, because 3-formylchromones without any halogen atom at the 8-position in the literature do not show the urease inhibitory activity (Kawase *et al.* 2007).

2. Experimental

Single crystals suitable for X-ray diffraction were obtained by slow evaporation of a 2-butanone solution of commercially available 6,8-dichloro-3-formylchromone at room temperature.

3. Refinement

The $C(sp^2)$ -bound hydrogen atoms were placed in geometrical positions [$C-H$ 0.95 Å, $U_{iso}(H) = 1.2U_{eq}(C)$], and refined using a riding model.

Computing details

Data collection: *WinAFC* (Rigaku, 1999); cell refinement: *WinAFC* (Rigaku, 1999); data reduction: *WinAFC* (Rigaku, 1999); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *CrystalStructure* (Rigaku, 2010); software used to prepare material for

publication: *CrystalStructure* (Rigaku, 2010).

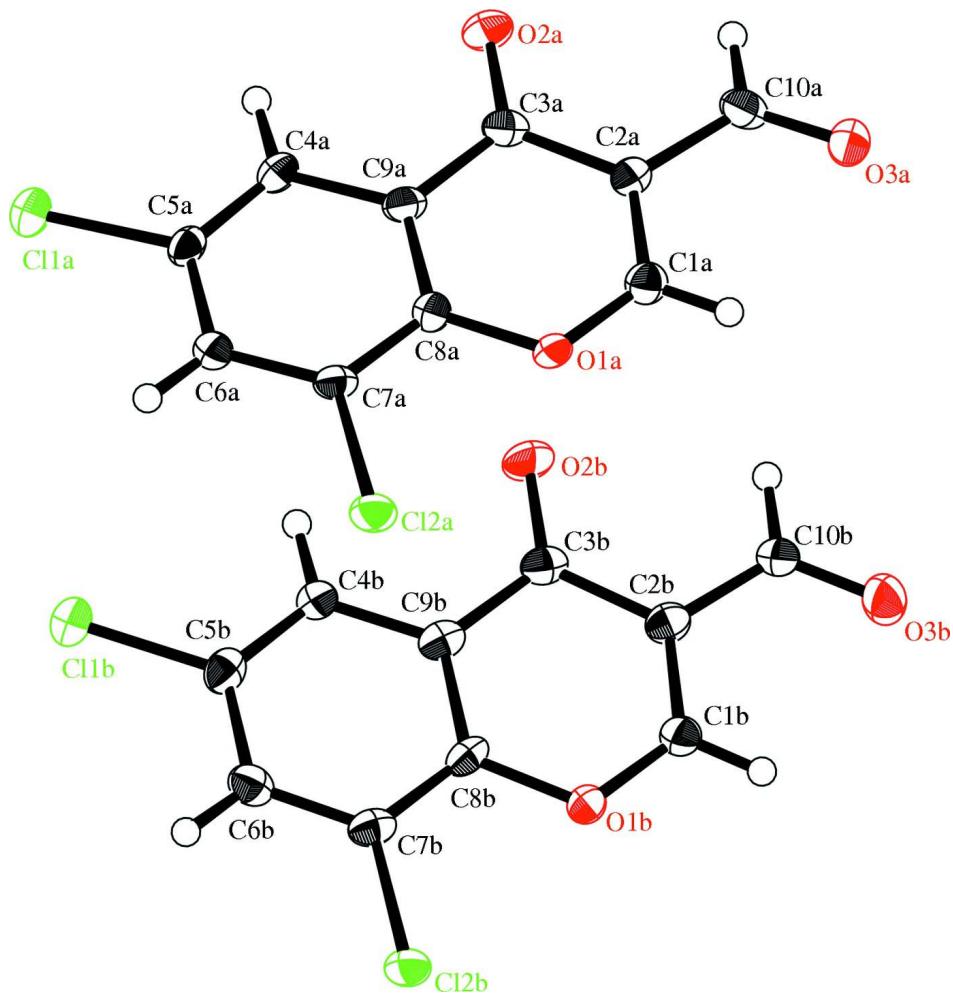
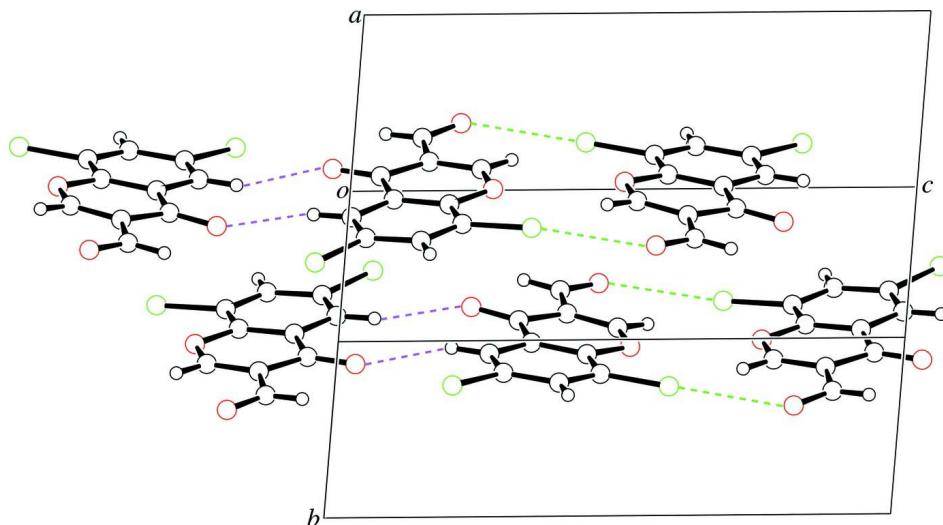


Figure 1

The molecular structure of the title compound, with displacement ellipsoids drawn at the 50% probability level. Hydrogen atoms are shown as small spheres of arbitrary radius.

**Figure 2**

A view of the intermolecular interactions of the title compound, represented as dashed green lines for $\text{Cl}\cdots\text{O}$ and dashed magenta lines for $\text{C}-\text{H}\cdots\text{O}$ interactions.

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 $b = 8.325$ (7) Å
 $c = 13.706$ (7) Å
 $\alpha = 96.55$ (6)°
 $\beta = 92.23$ (7)°
 $\gamma = 101.98$ (7)°
 $V = 917.2$ (13) Å³

$Z = 4$
 $F(000) = 488.00$
 $D_x = 1.760 \text{ Mg m}^{-3}$
Mo $K\alpha$ radiation, $\lambda = 0.71069$ Å
Cell parameters from 23 reflections
 $\theta = 15.2\text{--}17.4^\circ$
 $\mu = 0.68 \text{ mm}^{-1}$
 $T = 100$ K
Prismatic, colourless
 $0.42 \times 0.22 \times 0.08$ mm

Data collection

Rigaku AFC-7R
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 ω -2θ scans
Absorption correction: ψ scan
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2596 reflections with $F^2 > 2\sigma(F^2)$
 $R_{\text{int}} = 0.057$
 $\theta_{\text{max}} = 27.5^\circ$
 $h = -10 \rightarrow 6$
 $k = -10 \rightarrow 10$
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3 standard reflections every 150 reflections
intensity decay: 4.9%

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.076$
 $wR(F^2) = 0.212$
 $S = 1.10$
4203 reflections
271 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

$$w = 1/[\sigma^2(F_o^2) + (0.0531P)^2 + 6.620P]$$

$$\text{where } P = (F_o^2 + 2F_c^2)/3$$

$$(\Delta/\sigma)_{\max} < 0.001$$

$$\Delta\rho_{\max} = 0.70 \text{ e \AA}^{-3}$$

$$\Delta\rho_{\min} = -0.79 \text{ e \AA}^{-3}$$

Special details

Refinement. Refinement was performed using all reflections. The weighted *R*-factor (*wR*) and goodness of fit (*S*) are based on F^2 . *R*-factor (gt) are based on *F*. The threshold expression of $F^2 > 2.0 \sigma(F^2)$ is used only for calculating *R*-factor (gt).

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (Å²)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
Cl1a	0.9284 (2)	0.7180 (2)	-0.05206 (12)	0.0275 (4)
Cl2a	0.8997 (2)	0.60177 (19)	0.32529 (11)	0.0243 (4)
Cl1b	0.8654 (3)	1.0637 (2)	0.20783 (12)	0.0293 (4)
Cl2b	0.7665 (2)	1.01839 (19)	0.59010 (11)	0.0246 (4)
O1a	0.6255 (6)	0.33339 (5)	0.2521 (3)	0.0218 (10)
O2a	0.4506 (6)	0.1743 (6)	-0.0336 (4)	0.0277 (11)
O3a	0.2398 (7)	-0.0767 (6)	0.1869 (4)	0.0319 (12)
O1b	0.4940 (6)	0.7493 (6)	0.5162 (3)	0.0219 (10)
O2b	0.3559 (6)	0.5491 (6)	0.2305 (3)	0.0291 (11)
O3b	0.1021 (7)	0.3424 (6)	0.4522 (4)	0.0326 (12)
C1a	0.4980 (8)	0.2042 (8)	0.2288 (5)	0.0221 (13)
C2a	0.4342 (8)	0.1454 (8)	0.1358 (5)	0.0197 (13)
C3a	0.5023 (8)	0.2255 (8)	0.0520 (5)	0.0217 (13)
C4a	0.7126 (9)	0.4638 (8)	0.0063 (5)	0.0242 (14)
C5a	0.8382 (8)	0.5956 (8)	0.0344 (5)	0.0201 (13)
C6a	0.9020 (8)	0.6399 (8)	0.1321 (5)	0.0202 (13)
C7a	0.8290 (8)	0.5497 (8)	0.2026 (5)	0.0197 (13)
C8a	0.6945 (8)	0.4171 (8)	0.1770 (5)	0.0202 (13)
C9a	0.6370 (9)	0.3695 (8)	0.0784 (5)	0.0207 (13)
C10a	0.2954 (9)	-0.0026 (8)	0.1204 (5)	0.0250 (14)
C1b	0.3630 (8)	0.6197 (8)	0.4918 (5)	0.0220 (13)
C2b	0.3130 (8)	0.5479 (8)	0.4003 (5)	0.0217 (13)
C3b	0.3958 (8)	0.6127 (8)	0.3161 (5)	0.0214 (13)
C4b	0.6216 (8)	0.8361 (8)	0.2689 (5)	0.0220 (13)
C5b	0.7526 (8)	0.9654 (8)	0.2961 (5)	0.0227 (14)
C6b	0.8016 (9)	1.0234 (8)	0.3958 (5)	0.0234 (14)
C7b	0.7127 (8)	0.9495 (8)	0.4674 (5)	0.0217 (13)
C8b	0.5792 (8)	0.8170 (8)	0.4411 (5)	0.0188 (13)
C9b	0.5333 (8)	0.7557 (8)	0.3414 (5)	0.0190 (13)
C10b	0.1768 (9)	0.4034 (8)	0.3852 (5)	0.0238 (14)
H1a	0.4491	0.1499	0.2811	0.0265*
H2a	0.6755	0.4349	-0.0611	0.0290*
H3a	0.9933	0.7302	0.1493	0.0243*
H4a	0.2484	-0.0406	0.0555	0.0300*
H1b	0.3024	0.5769	0.5440	0.0264*
H2b	0.5904	0.8006	0.2012	0.0264*
H3b	0.8945	1.1120	0.4131	0.0281*
H4b	0.1440	0.3536	0.3196	0.0286*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Cl1a	0.0338 (9)	0.0277 (9)	0.0216 (8)	0.0044 (7)	0.0053 (7)	0.0088 (6)
Cl2a	0.0312 (9)	0.0231 (8)	0.0166 (7)	0.0037 (7)	-0.0028 (6)	-0.0007 (6)
Cl1b	0.0345 (10)	0.0287 (9)	0.0246 (8)	0.0034 (8)	0.0082 (7)	0.0071 (7)
Cl2b	0.0311 (9)	0.0232 (8)	0.0171 (7)	0.0031 (7)	-0.0008 (6)	-0.0016 (6)
O1a	0.034 (3)	0.019 (3)	0.012 (2)	0.0034 (19)	0.0043 (18)	0.0014 (16)
O2a	0.035 (3)	0.032 (3)	0.014 (3)	0.005 (3)	-0.0023 (19)	-0.0005 (18)
O3a	0.037 (3)	0.030 (3)	0.023 (3)	-0.005 (3)	0.006 (2)	0.002 (2)
O1b	0.029 (3)	0.021 (3)	0.014 (2)	0.0013 (19)	0.0014 (18)	0.0028 (17)
O2b	0.035 (3)	0.035 (3)	0.014 (3)	0.003 (3)	-0.0017 (19)	-0.0023 (19)
O3b	0.036 (3)	0.032 (3)	0.025 (3)	-0.007 (3)	-0.002 (3)	0.005 (2)
C1a	0.025 (4)	0.022 (4)	0.020 (3)	0.006 (3)	0.001 (3)	0.004 (3)
C2a	0.020 (4)	0.021 (3)	0.018 (3)	0.005 (3)	0.003 (3)	0.001 (3)
C3a	0.026 (4)	0.020 (3)	0.019 (3)	0.007 (3)	-0.001 (3)	-0.001 (3)
C4a	0.033 (4)	0.025 (4)	0.013 (3)	0.003 (3)	0.003 (3)	0.004 (3)
C5a	0.028 (4)	0.019 (3)	0.017 (3)	0.010 (3)	0.007 (3)	0.006 (3)
C6a	0.025 (4)	0.017 (3)	0.019 (3)	0.004 (3)	0.003 (3)	0.002 (3)
C7a	0.025 (4)	0.020 (3)	0.014 (3)	0.006 (3)	-0.001 (3)	-0.001 (3)
C8a	0.024 (4)	0.019 (3)	0.017 (3)	0.004 (3)	0.002 (3)	0.004 (3)
C9a	0.030 (4)	0.018 (3)	0.015 (3)	0.009 (3)	0.002 (3)	0.001 (3)
C10a	0.030 (4)	0.019 (3)	0.024 (4)	0.004 (3)	-0.002 (3)	-0.001 (3)
C1b	0.027 (4)	0.019 (3)	0.021 (3)	0.006 (3)	0.004 (3)	0.002 (3)
C2b	0.023 (4)	0.025 (4)	0.018 (3)	0.007 (3)	-0.001 (3)	0.002 (3)
C3b	0.028 (4)	0.022 (4)	0.017 (3)	0.011 (3)	-0.000 (3)	0.002 (3)
C4b	0.027 (4)	0.021 (3)	0.018 (3)	0.004 (3)	0.004 (3)	0.002 (3)
C5b	0.028 (4)	0.023 (4)	0.022 (4)	0.012 (3)	0.006 (3)	0.008 (3)
C6b	0.027 (4)	0.020 (4)	0.022 (4)	0.004 (3)	-0.003 (3)	0.001 (3)
C7b	0.026 (4)	0.024 (4)	0.016 (3)	0.010 (3)	-0.000 (3)	0.001 (3)
C8b	0.023 (4)	0.024 (4)	0.014 (3)	0.013 (3)	0.005 (3)	0.006 (3)
C9b	0.020 (3)	0.023 (3)	0.017 (3)	0.013 (3)	0.000 (3)	0.003 (3)
C10b	0.031 (4)	0.019 (3)	0.020 (3)	0.004 (3)	0.004 (3)	0.000 (3)

Geometric parameters (\AA , ^\circ)

Cl1a—C5a	1.740 (7)	C7a—C8a	1.398 (8)
Cl2a—C7a	1.735 (6)	C8a—C9a	1.401 (9)
Cl1b—C5b	1.733 (7)	C1b—C2b	1.338 (9)
Cl2b—C7b	1.723 (6)	C2b—C3b	1.464 (9)
O1a—C1a	1.344 (7)	C2b—C10b	1.458 (9)
O1a—C8a	1.381 (8)	C3b—C9b	1.465 (8)
O2a—C3a	1.229 (8)	C4b—C5b	1.367 (9)
O3a—C10a	1.210 (9)	C4b—C9b	1.405 (9)
O1b—C1b	1.363 (7)	C5b—C6b	1.411 (9)
O1b—C8b	1.379 (8)	C6b—C7b	1.376 (10)
O2b—C3b	1.233 (7)	C7b—C8b	1.393 (8)
O3b—C10b	1.227 (9)	C8b—C9b	1.412 (8)
C1a—C2a	1.356 (9)	C1a—H1a	0.950
C2a—C3a	1.467 (9)	C4a—H2a	0.950

C2a—C10a	1.490 (9)	C6a—H3a	0.950
C3a—C9a	1.459 (9)	C10a—H4a	0.950
C4a—C5a	1.355 (9)	C1b—H1b	0.950
C4a—C9a	1.416 (9)	C4b—H2b	0.950
C5a—C6a	1.401 (9)	C6b—H3b	0.950
C6a—C7a	1.373 (9)	C10b—H4b	0.950
C1a—O1a—C8a	118.3 (5)	C5b—C4b—C9b	119.9 (6)
C1b—O1b—C8b	118.1 (5)	C11b—C5b—C4b	120.6 (5)
O1a—C1a—C2a	124.6 (6)	C11b—C5b—C6b	117.6 (5)
C1a—C2a—C3a	120.4 (6)	C4b—C5b—C6b	121.9 (6)
C1a—C2a—C10a	118.9 (6)	C5b—C6b—C7b	118.9 (6)
C3a—C2a—C10a	120.7 (6)	C12b—C7b—C6b	120.4 (5)
O2a—C3a—C2a	122.6 (6)	C12b—C7b—C8b	119.6 (5)
O2a—C3a—C9a	122.9 (6)	C6b—C7b—C8b	120.0 (6)
C2a—C3a—C9a	114.5 (5)	O1b—C8b—C7b	117.3 (5)
C5a—C4a—C9a	119.4 (6)	O1b—C8b—C9b	121.6 (5)
C11a—C5a—C4a	120.3 (5)	C7b—C8b—C9b	121.1 (6)
C11a—C5a—C6a	117.0 (5)	C3b—C9b—C4b	121.9 (6)
C4a—C5a—C6a	122.7 (6)	C3b—C9b—C8b	119.8 (6)
C5a—C6a—C7a	118.3 (6)	C4b—C9b—C8b	118.2 (5)
C12a—C7a—C6a	120.3 (5)	O3b—C10b—C2b	123.8 (6)
C12a—C7a—C8a	119.0 (5)	O1a—C1a—H1a	117.700
C6a—C7a—C8a	120.6 (6)	C2a—C1a—H1a	117.702
O1a—C8a—C7a	117.4 (5)	C5a—C4a—H2a	120.314
O1a—C8a—C9a	122.2 (5)	C9a—C4a—H2a	120.307
C7a—C8a—C9a	120.4 (6)	C5a—C6a—H3a	120.850
C3a—C9a—C4a	121.6 (6)	C7a—C6a—H3a	120.844
C3a—C9a—C8a	119.9 (6)	O3a—C10a—H4a	118.481
C4a—C9a—C8a	118.5 (6)	C2a—C10a—H4a	118.489
O3a—C10a—C2a	123.0 (6)	O1b—C1b—H1b	117.382
O1b—C1b—C2b	125.2 (6)	C2b—C1b—H1b	117.374
C1b—C2b—C3b	120.2 (6)	C5b—C4b—H2b	120.070
C1b—C2b—C10b	119.4 (6)	C9b—C4b—H2b	120.068
C3b—C2b—C10b	120.3 (6)	C5b—C6b—H3b	120.568
O2b—C3b—C2b	122.6 (6)	C7b—C6b—H3b	120.567
O2b—C3b—C9b	122.5 (6)	O3b—C10b—H4b	118.104
C2b—C3b—C9b	114.9 (5)	C2b—C10b—H4b	118.102
C1a—O1a—C8a—C7a	-178.8 (6)	O1a—C8a—C9a—C3a	-1.4 (10)
C1a—O1a—C8a—C9a	-0.7 (9)	O1a—C8a—C9a—C4a	178.7 (6)
C8a—O1a—C1a—C2a	1.9 (10)	C7a—C8a—C9a—C3a	176.7 (6)
C8a—O1a—C1a—H1a	-178.1	C7a—C8a—C9a—C4a	-3.2 (10)
C1b—O1b—C8b—C7b	180.0 (6)	O1b—C1b—C2b—C3b	-2.9 (11)
C1b—O1b—C8b—C9b	0.3 (9)	O1b—C1b—C2b—C10b	176.3 (6)
C8b—O1b—C1b—C2b	2.5 (10)	H1b—C1b—C2b—C3b	177.1
C8b—O1b—C1b—H1b	-177.5	H1b—C1b—C2b—C10b	-3.7
O1a—C1a—C2a—C3a	-1.0 (11)	C1b—C2b—C3b—O2b	178.7 (7)
O1a—C1a—C2a—C10a	178.3 (6)	C1b—C2b—C3b—C9b	0.7 (10)

H1a—C1a—C2a—C3a	179.0	C1b—C2b—C10b—O3b	0.7 (11)
H1a—C1a—C2a—C10a	-1.7	C1b—C2b—C10b—H4b	-179.3
C1a—C2a—C3a—O2a	178.4 (7)	C3b—C2b—C10b—O3b	179.9 (7)
C1a—C2a—C3a—C9a	-1.1 (10)	C3b—C2b—C10b—H4b	-0.1
C1a—C2a—C10a—O3a	-2.4 (11)	C10b—C2b—C3b—O2b	-0.4 (11)
C1a—C2a—C10a—H4a	177.6	C10b—C2b—C3b—C9b	-178.5 (6)
C3a—C2a—C10a—O3a	176.9 (7)	O2b—C3b—C9b—C4b	4.2 (11)
C3a—C2a—C10a—H4a	-3.1	O2b—C3b—C9b—C8b	-176.2 (6)
C10a—C2a—C3a—O2a	-0.9 (11)	C2b—C3b—C9b—C4b	-177.7 (6)
C10a—C2a—C3a—C9a	179.7 (6)	C2b—C3b—C9b—C8b	1.8 (9)
O2a—C3a—C9a—C4a	2.6 (11)	C5b—C4b—C9b—C3b	-177.7 (6)
O2a—C3a—C9a—C8a	-177.3 (6)	C5b—C4b—C9b—C8b	2.7 (10)
C2a—C3a—C9a—C4a	-178.0 (6)	C9b—C4b—C5b—Cl1b	178.8 (6)
C2a—C3a—C9a—C8a	2.2 (10)	C9b—C4b—C5b—C6b	-0.9 (11)
C5a—C4a—C9a—C3a	-179.4 (6)	H2b—C4b—C5b—Cl1b	-1.2
C5a—C4a—C9a—C8a	0.5 (11)	H2b—C4b—C5b—C6b	179.1
C9a—C4a—C5a—Cl1a	-177.7 (6)	H2b—C4b—C9b—C3b	2.3
C9a—C4a—C5a—C6a	2.5 (11)	H2b—C4b—C9b—C8b	-177.3
H2a—C4a—C5a—Cl1a	2.3	Cl1b—C5b—C6b—C7b	179.0 (5)
H2a—C4a—C5a—C6a	-177.5	Cl1b—C5b—C6b—H3b	-1.0
H2a—C4a—C9a—C3a	0.6	C4b—C5b—C6b—C7b	-1.3 (11)
H2a—C4a—C9a—C8a	-179.5	C4b—C5b—C6b—H3b	178.6
Cl1a—C5a—C6a—C7a	177.5 (5)	C5b—C6b—C7b—Cl2b	-179.1 (6)
Cl1a—C5a—C6a—H3a	-2.5	C5b—C6b—C7b—C8b	1.6 (11)
C4a—C5a—C6a—C7a	-2.7 (11)	H3b—C6b—C7b—Cl2b	0.9
C4a—C5a—C6a—H3a	177.3	H3b—C6b—C7b—C8b	-178.4
C5a—C6a—C7a—Cl2a	-178.6 (6)	Cl2b—C7b—C8b—O1b	1.3 (9)
C5a—C6a—C7a—C8a	-0.1 (10)	Cl2b—C7b—C8b—C9b	-179.0 (5)
H3a—C6a—C7a—Cl2a	1.4	C6b—C7b—C8b—O1b	-179.3 (6)
H3a—C6a—C7a—C8a	179.9	C6b—C7b—C8b—C9b	0.3 (11)
Cl2a—C7a—C8a—O1a	-0.3 (9)	O1b—C8b—C9b—C3b	-2.4 (10)
Cl2a—C7a—C8a—C9a	-178.5 (5)	O1b—C8b—C9b—C4b	177.2 (6)
C6a—C7a—C8a—O1a	-178.8 (6)	C7b—C8b—C9b—C3b	178.0 (6)
C6a—C7a—C8a—C9a	3.0 (10)	C7b—C8b—C9b—C4b	-2.5 (10)

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

$D\text{—H}\cdots A$	$D\text{—H}$	$\text{H}\cdots A$	$D\cdots A$	$D\text{—H}\cdots A$
$\text{C}4b^{\text{i}}\text{—H}2b^{\text{i}}\cdots \text{O}2a$	0.95	2.35	3.246 (8)	157
$\text{C}4a\text{—H}2a\cdots \text{O}2b^{\text{i}}$	0.95	2.35	3.259 (8)	160

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