

Acta Crystallographica Section E

Structure Reports

Online

ISSN 1600-5368

2,4-Dibromo-1,3-dihydroxy-9H-xanthen-9-one

 Shi-Wen Huang,^{a,b} Zheng-Min Yang,^a Fu-Ping Huang^a and Jiang-Ke Qin^{a*}
^aKey Laboratory for the Chemistry & Molecular Engineering of Medicinal Resources, Ministry of Education of China, School of Chemistry & Chemical Engineering, Guangxi Normal University, Guilin, 541004, People's Republic of China, and

^bDepartment of Pharmacy, Youjiang Medical University for Nationalities, Baise 533000, People's Republic of China

Correspondence e-mail: jiangkeq@163.com

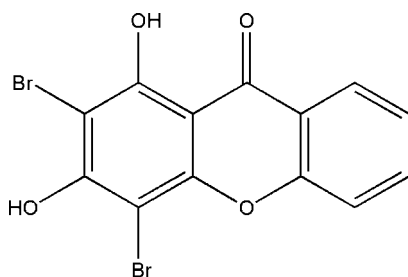
Received 26 May 2013; accepted 12 July 2013

 Key indicators: single-crystal X-ray study; $T = 298$ K; mean $\sigma(\text{C}-\text{C}) = 0.008$ Å; R factor = 0.034; wR factor = 0.070; data-to-parameter ratio = 12.3.

The title compound, $\text{C}_{13}\text{H}_6\text{Br}_2\text{O}_4$, derived from xanthone, a fundamental structural framework of active ingredients in many medicinal plants, and was synthesized by bromination of 1,3-dihydroxyxanthen-9-one with *N*-bromosuccinimide. The molecular conformation is essentially planar, the dihedral angle between the benzene rings being $1.1(4)^\circ$. This conformation is favorable for the formation of an intramolecular $\text{O}-\text{H}\cdots\text{O}$ hydrogen bond between a hydroxy group and the xanthone carbonyl group. In the crystal, molecules are associated into chains along the *b*-axis direction via $\text{C}=\text{O}\cdots\text{H}-\text{O}$ hydrogen bonds involving the other hydroxy group.

Related literature

For the pharmacological activity of xanthone derivatives, see: Cheng *et al.* (2011); Dao *et al.* (2012); Sousa *et al.* (2009); Szkaradek *et al.* (2013). For the synthesis of the xanthone used as a starting material, see: Liu *et al.* (2006). For related xanthone structures, see: Corrêa *et al.* (2010).



Experimental

Crystal data

 $\text{C}_{13}\text{H}_6\text{Br}_2\text{O}_4$
 $M_r = 386.00$

 Orthorhombic, $Pna2_1$
 $a = 18.4489(15)$ Å
 $b = 16.9049(13)$ Å
 $c = 3.8564(3)$ Å
 $V = 1202.72(16)$ Å³
 $Z = 4$
 Mo $K\alpha$ radiation
 $\mu = 6.75$ mm⁻¹
 $T = 298$ K
 $0.28 \times 0.09 \times 0.06$ mm

Data collection

 Bruker SMART CCD diffractometer
 Absorption correction: multi-scan (SADABS; Bruker, 1998)
 $T_{\min} = 0.254$, $T_{\max} = 0.688$

 6188 measured reflections
 2120 independent reflections
 1830 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.076$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.034$
 $wR(F^2) = 0.070$
 $S = 1.04$
 2120 reflections
 172 parameters
 1 restraint
 H-atom parameters constrained

 $\Delta\rho_{\max} = 0.43$ e Å⁻³
 $\Delta\rho_{\min} = -0.33$ e Å⁻³
 Absolute structure: Flack (1983),
 881 Friedel pairs
 Absolute structure parameter:
 $-0.008(16)$
Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{O4}-\text{H4}\cdots\text{O2}$	0.82	1.81	2.555 (7)	149
$\text{O3}-\text{H3}\cdots\text{O2}^i$	0.82	2.02	2.741 (7)	147

 Symmetry code: (i) $-x + \frac{3}{2}, y + \frac{1}{2}, z - \frac{1}{2}$.

Data collection: SMART (Bruker, 1998); cell refinement: SAINT (Bruker, 1998); data reduction: SAINT (Bruker, 1998); program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: SHELXL97 (Sheldrick, 2008); software used to prepare material for publication: SHELXL97 (Sheldrick, 2008).

This work was supported financially by grants from the National Natural Science Foundation of PRC (21002015) and the Natural Science Foundation of Guangxi (2010GXNSFB013013).

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: BH2480).

References

- Bruker (1998). SMART, SAINT and SADABS. Bruker AXS Inc., Madison, Wisconsin, USA.
- Cheng, J.-H., Huang, A.-M., Hour, T.-C., Yang, S.-C., Pu, Y.-S. & Lin, C.-N. (2011). *Eur. J. Med. Chem.* **46**, 1222–1231.
- Corrêa, R. S., dos Santos, M. H., Nagem, T. J. & Ellena, J. (2010). *Struct. Chem.* **21**, 555–563.
- Dao, T. T., Dang, T. T., Nguyen, P. H., Kim, E., Thuong, P. T. & Oh, W. K. (2012). *Bioorg. Med. Chem. Lett.* **22**, 3688–3692.
- Flack, H. D. (1983). *Acta Cryst.* **A39**, 876–881.
- Liu, Y., Zou, L., Ma, L., Chen, W.-H., Wang, B. & Xu, Z.-L. (2006). *Bioorg. Med. Chem.* **14**, 5683–5690.
- Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112–122.
- Sousa, E., Paiva, A., Nazareth, N., Gales, L., Damas, A. M., Nascimento, M. S. J. & Pinto, M. (2009). *Eur. J. Med. Chem.* **44**, 3830–3835.
- Szkaradek, N., Gunia, A., Waszkielewicz, A. M., Antkiewicz-Michaluk, L., Cegla, M., Szneler, E. & Marona, H. (2013). *Bioorg. Med. Chem.* **21**, 1190–1198.

supplementary materials

Acta Cryst. (2013). E69, o1361 [doi:10.1107/S1600536813019296]

2,4-Dibromo-1,3-dihydroxy-9H-xanthen-9-one

Shi-Wen Huang, Zheng-Min Yang, Fu-Ping Huang and Jiang-Ke Qin

Comment

Xanthone, also named as dibenzo- γ -pyrone, is a fundamental structural framework of active ingredients in many medicinal plants, which derivatives have broad pharmacological activities, such as antioxidant (Cheng *et al.*, 2011), antitumor (Sousa *et al.*, 2009), anticonvulsant (Szkaradek *et al.*, 2013) and inhibition of neuraminidase activity (Dao *et al.*, 2012). The title compound in this study is a new xanthone derivative, which was synthesized by bromination of 1,3-dihydroxy-xanthen-9-one (Liu *et al.*, 2006) with NBS.

We report here the synthesis and crystal structure of 2,4-dibromo-1,3-dihydroxy-xanthen-9-one (C₁₃H₆Br₂O₄, Fig. 1). The molecule of the title compound has a planar conformation, with characteristic bond lengths C2—Br1 = 1.893 (5) Å and C4—Br2 = 1.902 (5) Å. The molecular conformation is mainly controlled by the O4—H4...O2 intramolecular hydrogen bond between the hydroxy OH group and the carbonyl O atom. Molecules are further connected into a one-dimensional supramolecular architecture *via* O3—H3...O2 intermolecular hydrogen bonds (Fig. 2). The title compound has the same planar molecular conformation as that reported for other 1-hydroxy-9H-xanthen-9-one derivatives (Corrêa *et al.*, 2010), while the carbonyl bond length C7=O2, 1.262 (7) Å, is slightly larger than the corresponding carbonyl bond lengths in these derivatives.

Experimental

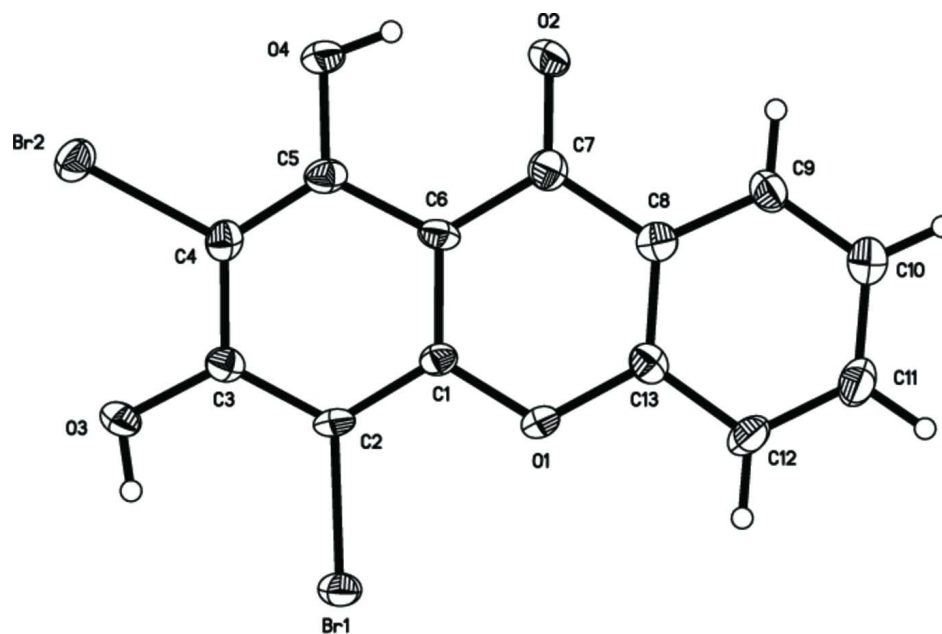
The title compound was synthesized using the following procedure: in a 50 ml flask, 1,3-dihydroxy-xanthen-9-one (1010 mg, 4.43 mmol; Liu *et al.*, 2006) was dissolved in CCl₄ (15 ml), then NBS (500 mg, 2.83 mmol) was added. The mixture was stirred at room temperature for 24 h. The residue was washed with acetone and then filtered. The yellow solid was collected and dried. Recrystallization from methanol solution afforded 2,4-dibromo-1,3-dihydroxy-xanthen-9-one as yellow crystals. The compound identity was confirmed by NMR spectroscopy. ¹H NMR (500 MHz, DMSO-*d*₆): 13.63 (*s*, 1H), 8.09 (*dd*, *J* = 7.9, 1.6 Hz, 1H), 7.87 (*td*, *J* = 7.8, 1.5 Hz, 1H), 7.59 (*d*, *J* = 8.4 Hz, 1H), 7.48 (*t*, *J* = 7.5 Hz, 1H).

Refinement

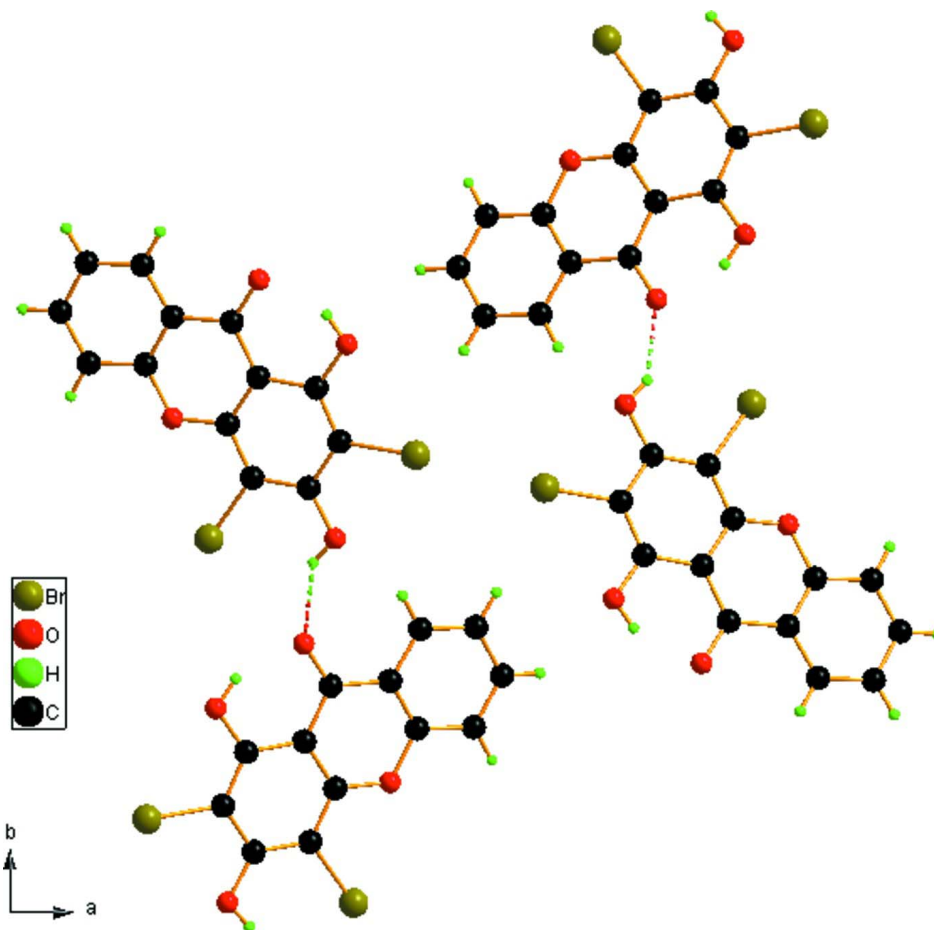
The H atoms on C and O atoms were positioned geometrically and refined using a riding model, with C—H = 0.93 Å and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$, and with O—H = 0.82 Å and $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{O})$.

Computing details

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINT* (Bruker, 1998); data reduction: *SAINT* (Bruker, 1998); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXL97* (Sheldrick, 2008); software used to prepare material for publication: *SHELXL97* (Sheldrick, 2008).

**Figure 1**

The structure of the title compound and the atom-numbering scheme.


Figure 2

The packing diagram of the title compound.

2,4-Dibromo-1,3-dihydroxy-9H-xanthen-9-one

Crystal data

$C_{13}H_6Br_2O_4$

$M_r = 386.00$

Orthorhombic, $Pna2_1$

Hall symbol: $P\ 2c\ -2n$

$a = 18.4489\ (15)\ \text{\AA}$

$b = 16.9049\ (13)\ \text{\AA}$

$c = 3.8564\ (3)\ \text{\AA}$

$V = 1202.72\ (16)\ \text{\AA}^3$

$Z = 4$

$F(000) = 744$

$D_x = 2.132\ \text{Mg m}^{-3}$

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

Cell parameters from 2778 reflections

$\theta = 3.3\text{--}26.2^\circ$

$\mu = 6.75\ \text{mm}^{-1}$

$T = 298\ \text{K}$

Block, yellow

$0.28 \times 0.09 \times 0.06\ \text{mm}$

Data collection

Bruker SMART CCD

diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

φ and ω scans

Absorption correction: multi-scan

(*SADABS*; Bruker, 1998)

$T_{\min} = 0.254$, $T_{\max} = 0.688$

6188 measured reflections

2120 independent reflections

1830 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.076$
 $\theta_{\text{max}} = 25.0^\circ$, $\theta_{\text{min}} = 1.6^\circ$
 $h = -21 \rightarrow 21$

$k = -20 \rightarrow 15$
 $l = -4 \rightarrow 4$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.034$
 $wR(F^2) = 0.070$
 $S = 1.04$
 2120 reflections
 172 parameters
 1 restraint
 0 constraints
 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
 $w = 1/[\sigma^2(F_o^2) + (0.0232P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.002$
 $\Delta\rho_{\text{max}} = 0.43 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.33 \text{ e } \text{\AA}^{-3}$
 Absolute structure: Flack (1983), 881 Friedel
 pairs
 Flack parameter: $-0.008 (16)$

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
Br1	0.65912 (3)	1.09241 (3)	-0.07056 (16)	0.03711 (16)
Br2	0.91885 (3)	0.97740 (3)	0.54899 (17)	0.03972 (17)
O1	0.61335 (17)	0.9267 (2)	0.0466 (12)	0.0366 (9)
O2	0.72160 (18)	0.7368 (2)	0.4658 (13)	0.0483 (11)
O3	0.8177 (2)	1.0920 (2)	0.2248 (11)	0.0408 (11)
H3	0.7904	1.1248	0.1377	0.061*
O4	0.83232 (18)	0.8253 (2)	0.5513 (13)	0.0452 (11)
H4	0.8074	0.7851	0.5521	0.068*
C1	0.6824 (3)	0.9348 (3)	0.1582 (13)	0.0262 (13)
C2	0.7137 (3)	1.0086 (3)	0.1241 (13)	0.0277 (13)
C3	0.7844 (3)	1.0223 (3)	0.2401 (14)	0.0293 (13)
C4	0.8233 (3)	0.9589 (3)	0.3828 (15)	0.0310 (14)
C5	0.7941 (3)	0.8853 (3)	0.4141 (16)	0.0323 (13)
C6	0.7210 (3)	0.8704 (3)	0.2998 (14)	0.0305 (14)
C7	0.6879 (3)	0.7943 (4)	0.3340 (15)	0.0351 (15)
C8	0.6131 (3)	0.7875 (3)	0.2015 (14)	0.0337 (14)
C9	0.5753 (3)	0.7164 (4)	0.2048 (17)	0.0400 (15)
H9	0.5977	0.6707	0.2858	0.048*
C10	0.5053 (3)	0.7135 (4)	0.0887 (19)	0.0479 (16)
H10	0.4797	0.6661	0.0978	0.057*
C11	0.4724 (3)	0.7802 (4)	-0.0412 (18)	0.0458 (15)
H11	0.4251	0.7770	-0.1236	0.055*
C12	0.5080 (3)	0.8511 (3)	-0.0515 (17)	0.0410 (14)
H12	0.4853	0.8965	-0.1333	0.049*
C13	0.5793 (3)	0.8533 (3)	0.0642 (17)	0.0314 (12)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Br1	0.0367 (3)	0.0309 (3)	0.0438 (3)	0.0061 (2)	-0.0024 (3)	0.0080 (3)

Br2	0.0277 (3)	0.0461 (4)	0.0454 (3)	-0.0007 (2)	-0.0040 (3)	0.0016 (3)
O1	0.0249 (19)	0.033 (2)	0.052 (2)	0.0044 (15)	-0.007 (2)	0.008 (2)
O2	0.037 (2)	0.027 (2)	0.081 (3)	0.0009 (17)	-0.006 (3)	0.015 (2)
O3	0.040 (2)	0.025 (2)	0.058 (3)	-0.0020 (19)	-0.005 (2)	0.012 (2)
O4	0.032 (2)	0.037 (2)	0.067 (3)	0.0062 (16)	-0.010 (3)	0.015 (2)
C1	0.025 (3)	0.026 (3)	0.027 (3)	0.006 (2)	0.000 (2)	-0.002 (2)
C2	0.028 (3)	0.028 (3)	0.026 (4)	0.012 (2)	-0.002 (2)	0.001 (2)
C3	0.028 (3)	0.028 (3)	0.032 (3)	0.003 (3)	0.005 (2)	0.003 (3)
C4	0.022 (3)	0.036 (3)	0.034 (3)	0.000 (2)	0.006 (3)	-0.002 (3)
C5	0.027 (3)	0.036 (3)	0.034 (3)	0.006 (2)	0.005 (3)	0.008 (3)
C6	0.033 (3)	0.023 (3)	0.036 (3)	0.008 (2)	0.003 (2)	0.005 (2)
C7	0.028 (3)	0.037 (4)	0.041 (4)	0.001 (3)	0.008 (3)	0.003 (3)
C8	0.037 (4)	0.034 (4)	0.031 (3)	0.002 (3)	-0.001 (3)	-0.003 (3)
C9	0.038 (4)	0.028 (3)	0.054 (4)	-0.004 (3)	0.002 (3)	0.001 (3)
C10	0.040 (4)	0.048 (4)	0.056 (4)	-0.008 (3)	-0.003 (3)	0.003 (3)
C11	0.030 (3)	0.056 (4)	0.051 (4)	-0.007 (3)	-0.004 (3)	0.001 (4)
C12	0.028 (3)	0.048 (4)	0.048 (4)	0.006 (3)	-0.008 (3)	-0.001 (3)
C13	0.034 (3)	0.032 (3)	0.029 (3)	-0.001 (2)	0.010 (3)	0.001 (3)

Geometric parameters (Å, °)

Br1—C2	1.893 (5)	C5—C6	1.442 (7)
Br2—C4	1.902 (5)	C6—C7	1.429 (8)
O1—C1	1.351 (6)	C7—C8	1.476 (8)
O1—C13	1.393 (6)	C8—C13	1.379 (7)
O2—C7	1.262 (7)	C8—C9	1.390 (8)
O3—C3	1.331 (7)	C9—C10	1.368 (8)
O3—H3	0.8200	C9—H9	0.9300
O4—C5	1.343 (6)	C10—C11	1.374 (8)
O4—H4	0.8200	C10—H10	0.9300
C1—C2	1.381 (7)	C11—C12	1.368 (8)
C1—C6	1.411 (7)	C11—H11	0.9300
C2—C3	1.398 (7)	C12—C13	1.389 (7)
C3—C4	1.402 (7)	C12—H12	0.9300
C4—C5	1.361 (7)		
C1—O1—C13	119.9 (4)	O2—C7—C6	121.3 (5)
C3—O3—H3	109.5	O2—C7—C8	122.7 (5)
C5—O4—H4	109.5	C6—C7—C8	115.9 (5)
O1—C1—C2	117.1 (4)	C13—C8—C9	118.4 (5)
O1—C1—C6	121.4 (5)	C13—C8—C7	119.5 (5)
C2—C1—C6	121.5 (5)	C9—C8—C7	122.2 (5)
C1—C2—C3	120.6 (5)	C10—C9—C8	120.0 (6)
C1—C2—Br1	119.4 (4)	C10—C9—H9	120.0
C3—C2—Br1	120.0 (4)	C8—C9—H9	120.0
O3—C3—C2	124.2 (5)	C9—C10—C11	120.5 (6)
O3—C3—C4	117.3 (5)	C9—C10—H10	119.8
C2—C3—C4	118.5 (5)	C11—C10—H10	119.8
C5—C4—C3	122.1 (5)	C12—C11—C10	121.2 (5)
C5—C4—Br2	119.1 (4)	C12—C11—H11	119.4

C3—C4—Br2	118.8 (4)	C10—C11—H11	119.4
O4—C5—C4	121.2 (5)	C11—C12—C13	117.9 (5)
O4—C5—C6	118.6 (5)	C11—C12—H12	121.1
C4—C5—C6	120.2 (5)	C13—C12—H12	121.1
C1—C6—C7	121.0 (5)	C8—C13—C12	122.0 (5)
C1—C6—C5	117.1 (5)	C8—C13—O1	122.3 (5)
C7—C6—C5	121.9 (5)	C12—C13—O1	115.7 (5)
C13—O1—C1—C2	178.1 (5)	O4—C5—C6—C7	1.1 (9)
C13—O1—C1—C6	-1.2 (8)	C4—C5—C6—C7	-178.8 (5)
O1—C1—C2—C3	178.6 (5)	C1—C6—C7—O2	-178.7 (5)
C6—C1—C2—C3	-2.1 (8)	C5—C6—C7—O2	-0.1 (9)
O1—C1—C2—Br1	0.3 (6)	C1—C6—C7—C8	2.1 (8)
C6—C1—C2—Br1	179.6 (4)	C5—C6—C7—C8	-179.2 (5)
C1—C2—C3—O3	-178.2 (5)	O2—C7—C8—C13	179.3 (6)
Br1—C2—C3—O3	0.1 (8)	C6—C7—C8—C13	-1.6 (8)
C1—C2—C3—C4	1.7 (8)	O2—C7—C8—C9	-1.8 (9)
Br1—C2—C3—C4	-180.0 (4)	C6—C7—C8—C9	177.3 (5)
O3—C3—C4—C5	179.3 (5)	C13—C8—C9—C10	-3.0 (9)
C2—C3—C4—C5	-0.6 (9)	C7—C8—C9—C10	178.0 (6)
O3—C3—C4—Br2	1.0 (7)	C8—C9—C10—C11	1.9 (11)
C2—C3—C4—Br2	-178.9 (4)	C9—C10—C11—C12	-1.3 (12)
C3—C4—C5—O4	179.9 (6)	C10—C11—C12—C13	1.9 (10)
Br2—C4—C5—O4	-1.8 (8)	C9—C8—C13—C12	3.7 (9)
C3—C4—C5—C6	-0.2 (9)	C7—C8—C13—C12	-177.4 (6)
Br2—C4—C5—C6	178.1 (4)	C9—C8—C13—O1	-179.2 (6)
O1—C1—C6—C7	-0.8 (8)	C7—C8—C13—O1	-0.3 (9)
C2—C1—C6—C7	179.9 (5)	C11—C12—C13—C8	-3.1 (10)
O1—C1—C6—C5	-179.5 (5)	C11—C12—C13—O1	179.6 (6)
C2—C1—C6—C5	1.2 (8)	C1—O1—C13—C8	1.7 (9)
O4—C5—C6—C1	179.8 (5)	C1—O1—C13—C12	179.0 (5)
C4—C5—C6—C1	-0.1 (8)		

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

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
O3—H3 \cdots Br1	0.82	2.61	3.139 (5)	124
O4—H4 \cdots O2	0.82	1.81	2.555 (7)	149
O3—H3 \cdots O2 ⁱ	0.82	2.02	2.741 (7)	147

Symmetry code: (i) $-x+3/2, y+1/2, z-1/2$.