

Acta Crystallographica Section E

Structure Reports

Online

ISSN 1600-5368

Ethyl 4-(5-bromo-2-hydroxyphenyl)-2,7,7-trimethyl-5-oxo-1,4,5,6,7,8-hexahydroquinoline-3-carboxylate

 Malahat M. Kurbanova,^a Elnur Z. Huseynov,^a Atash V. Gurbanov,^{a*} Abel M. Maharramov^a and Reza Kia^{b,c}
^aDepartment of Organic Chemistry, Baku State University, Baku, Azerbaijan,

^bDepartment of Chemistry, Science and Research Branch, Islamic Azad University, Tehran, Iran, and ^cStructural Dynamics of (Bio)Chemical Systems, Max Planck Institute for Biophysical Chemistry, Am Fassberg 11, 37077 Göttingen, Germany
Correspondence e-mail: organik10@hotmail.com

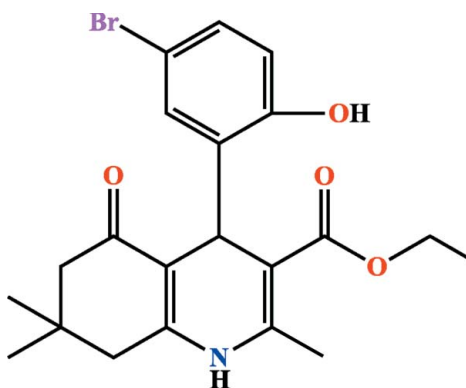
Received 4 March 2013; accepted 8 March 2013

 Key indicators: single-crystal X-ray study; $T = 294$ K; mean $\sigma(\text{C}-\text{C}) = 0.004$ Å; disorder in main residue; R factor = 0.050; wR factor = 0.136; data-to-parameter ratio = 18.8.

In the title compound, $\text{C}_{21}\text{H}_{24}\text{BrNO}_4$, the dihedral angle between the heterocyclic ring and the pendant aromatic ring is $80.20(13)^\circ$. The hexahydroquinone [*i.e.* the one with the $\text{C}=\text{O}$ group] ring adopts a sofa conformation. An intramolecular $\text{O}-\text{H}\cdots\text{O}$ hydrogen bond generates an $S(6)$ ring motif. The ethyl group is disordered over two sets of sites with a refined site occupancy ratio of 0.633 (10):0.366 (10). In the crystal, molecules are linked by $\text{N}-\text{H}\cdots\text{O}$ interactions, forming chains parallel to $[101]$. There are no significant $\text{C}-\text{H}\cdots\pi$ or $\pi-\pi$ interactions in the crystal structure.

Related literature

For standard bond lengths, see: Allen *et al.* (1987). For hydrogen-bond motifs, see: Bernstein *et al.* (1995). For background to hexahydroquinoline compounds and their applications, see: Sausins & Duburs (1988); Nakayama & Kasoaka (1996); Klusa (1995). For the synthesis of related compounds, see: Kumar *et al.* (2008); Song *et al.* (2012).



Experimental

Crystal data

$\text{C}_{21}\text{H}_{24}\text{BrNO}_4$	$V = 2009.84(10) \text{ \AA}^3$
$M_r = 434.32$	$Z = 4$
Monoclinic, $P2_1/n$	Mo $K\alpha$ radiation
$a = 9.5969(3) \text{ \AA}$	$\mu = 2.07 \text{ mm}^{-1}$
$b = 19.0805(5) \text{ \AA}$	$T = 294 \text{ K}$
$c = 11.0678(3) \text{ \AA}$	$0.24 \times 0.22 \times 0.18 \text{ mm}$
$\beta = 97.387(1)^\circ$	

Data collection

Bruker SMART APEXII CCD area-detector diffractometer	23241 measured reflections
Absorption correction: multi-scan (SADABS; Bruker, 2005)	5008 independent reflections
$T_{\min} = 0.636$, $T_{\max} = 0.707$	3604 reflections with $I > 2\sigma(I)$
	$R_{\text{int}} = 0.022$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.050$	3 restraints
$wR(F^2) = 0.136$	H-atom parameters constrained
$S = 1.05$	$\Delta\rho_{\text{max}} = 1.10 \text{ e \AA}^{-3}$
5008 reflections	$\Delta\rho_{\text{min}} = -1.02 \text{ e \AA}^{-3}$
266 parameters	

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{O4}-\text{H4}\cdots\text{O1}$	0.88	1.75	2.625 (3)	171
$\text{N1}-\text{H1}\cdots\text{O2}^i$	0.86	2.05	2.866 (3)	158

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

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, 2009).

The authors thank the Chemistry Department, Baku State University, for providing the X-ray diffraction facility.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: RZ5051).

References

- Allen, F. H., Kennard, O., Watson, D. G., Brammer, L., Orpen, A. G. & Taylor, R. (1987). *J. Chem. Soc. Perkin Trans. 2*, pp. S1–19.
- Bernstein, J., Davis, R. E., Shimoni, L. & Chang, N.-L. (1995). *Angew. Chem. Int. Ed. Engl.* **34**, 1555–1573.
- Bruker (2005). APEX2, SAINT and SADABS. Bruker AXS Inc., Madison, Wisconsin, USA.
- Klusa, V. (1995). *Drugs Future*, **20**, 135–138.
- Kumar, S., Sharma, P., Kapoor, K. K. & Hundal, M. S. (2008). *Tetrahedron*, **64**, 536–542.
- Nakayama, H. & Kasoaka, Y. (1996). *Heterocycles*, **42**, 901–909.
- Sausins, A. & Duburs, G. (1988). *Heterocycles*, **27**, 269–289.
- Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112–122.
- Song, S. J., Shan, Z. X. & Jin, J. (2012). *Synth. Commun.* **40**, 3067–3077.
- Spek, A. L. (2009). *Acta Cryst.* **D65**, 148–155.

supplementary materials

Acta Cryst. (2013). E69, o541 [doi:10.1107/S1600536813006739]

Ethyl 4-(5-bromo-2-hydroxyphenyl)-2,7,7-trimethyl-5-oxo-1,4,5,6,7,8-hexahydroquinoline-3-carboxylate

Malahat M. Kurbanova, Elnur Z. Huseynov, Atash V. Gurbanov, Abel M. Maharramov and Reza Kia

Comment

Hexahydroquinoline derivatives possess a variety of biological activities, such as vasodilatory, bronchodilatory, antiatherosclerotic, hepatoprotective, and antidiabetic activity (Sausins *et al.*, 1988), and some of them have been used as calcium channel modulators and curatives for cardiovascular diseases (Nakayama *et al.*, 1996). In past years, their uses as neuroprotectants, platelet anti-aggregatory agents, and cerebral anti-ischemic agents in the treatment of Alzheimer's disease and as chemosensitizers in tumor therapy have been also reported (Klusa, 1995).

The asymmetric unit of the title compound, Fig. 1, comprises a substituted hexahydroquinoline compound. Both six-membered rings of the hexahydroquinoline ring system adopt a half-boat conformation. Bond lengths (Allen *et al.*, 1987) and angles are within normal ranges. An intramolecular O—H \cdots O hydrogen bond generates an *S(6)* ring motif (Bernstein *et al.*, 1995). In the crystal structure, molecules are linked together by intermolecular N—H \cdots O hydrogen interactions (Table 1, Fig. 2) forming chains parallel to the [101] direction. The ethyl group is disordered over two sets of sites with a refined site occupancy ratio of 0.633 (10):0.366 (10). The compound contains one chiral center but the space group is centrosymmetric, so the molecule exists as a racemate.

Experimental

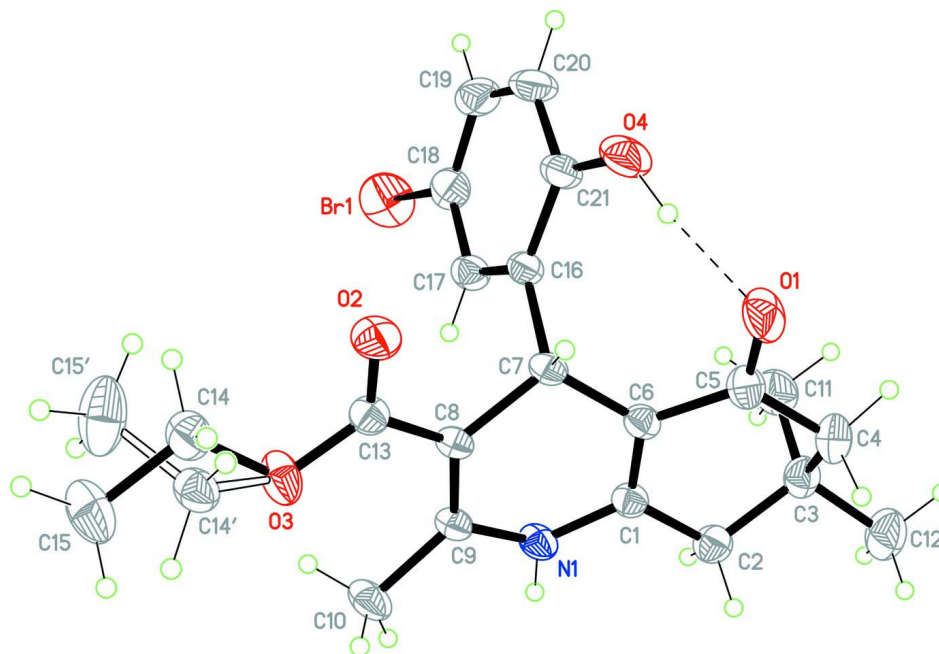
5-Bromsalicylaldehyde (0.201 g, 1 mmol), ethyl acetoacetate (0.25 ml, 1 mmol), dimedone (0.14 g, 1 mmol), ammonium acetate (0.116 g, 1.5 mmol) and ethanol (15 ml) were charged in a round bottom flask. Then the reaction mixture was stirred at room temperature for 12 hours, then the product was separated by filtration. Recrystallization was effected by using ethanol as solvent. Yield 86%. M. p. 520 K.

Refinement

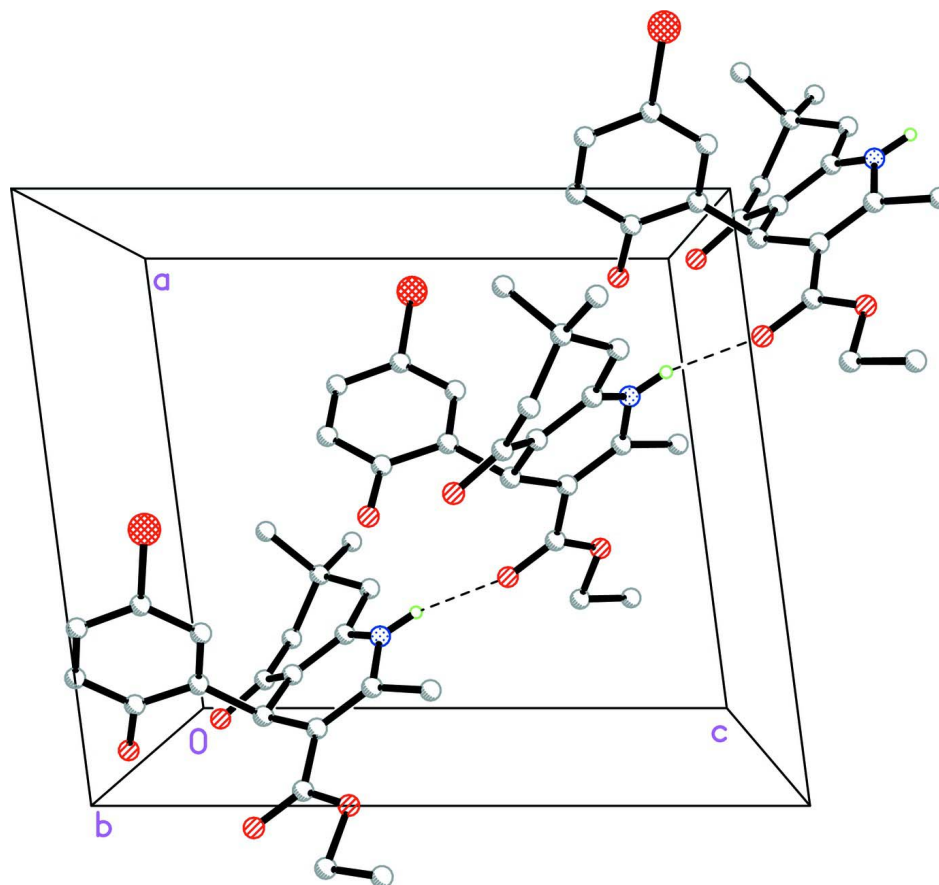
The O- and N-bound H atoms were located in a difference Fourier map and constrained to ride on their parent atoms with O—H = 0.88 Å, N—H = 0.86 Å and $U_{\text{iso}}(\text{H}) = 1.5 U_{\text{eq}}(\text{O}, \text{N})$. The C-bound H-atoms were included in calculated positions and treated as riding atoms with C—H = 0.93–0.98 Å and $U_{\text{iso}}(\text{H}) = 1.2 U_{\text{eq}}(\text{C})$ or $1.5 U_{\text{eq}}(\text{C})$ for methyl H atoms. Distance restraints were applied to the components of the disordered ethyl group.

Computing details

Data collection: *APEX2* (Bruker, 2005); cell refinement: *SAINT* (Bruker, 2005); data reduction: *SAINT* (Bruker, 2005); program(s) used to solve structure: *SHELXTL* (Sheldrick, 2008); program(s) used to refine structure: *SHELXTL* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL* (Sheldrick, 2008) and *PLATON* (Spek, 2009).

**Figure 1**

The molecular structure of the title compound, showing 40% probability displacement ellipsoids.

**Figure 2**

Packing diagram of the title compound viewed down the *b* axis, showing the linkage of molecules through N—H···O hydrogen interactions (dashed lines). Only H atoms involved in hydrogen bonding are shown.

Ethyl 4-(5-bromo-2-hydroxyphenyl)-2,7,7-trimethyl-5-oxo-1,4,5,6,7,8-hexahydroquinoline-3-carboxylate

Crystal data

$C_{21}H_{24}BrNO_4$

$M_r = 434.32$

Monoclinic, $P2_1/n$

Hall symbol: $-P\ 2_1/n$

$a = 9.5969\ (3)\ \text{\AA}$

$b = 19.0805\ (5)\ \text{\AA}$

$c = 11.0678\ (3)\ \text{\AA}$

$\beta = 97.387\ (1)^\circ$

$V = 2009.84\ (10)\ \text{\AA}^3$

$Z = 4$

$F(000) = 896$

$D_x = 1.435\ \text{Mg m}^{-3}$

Melting point: 520 K

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

Cell parameters from 2045 reflections

$\theta = 3.3\text{--}27.5^\circ$

$\mu = 2.07\ \text{mm}^{-1}$

$T = 294\ \text{K}$

Block, colourless

$0.24 \times 0.22 \times 0.18\ \text{mm}$

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)

$T_{\min} = 0.636$, $T_{\max} = 0.707$

23241 measured reflections

5008 independent reflections

3604 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.022$
 $\theta_{\text{max}} = 28.3^\circ$, $\theta_{\text{min}} = 2.1^\circ$
 $h = -12 \rightarrow 12$

$k = -25 \rightarrow 25$
 $l = -14 \rightarrow 14$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.050$
 $wR(F^2) = 0.136$
 $S = 1.05$
 5008 reflections
 266 parameters
 3 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
 $w = 1/[\sigma^2(F_o^2) + (0.0542P)^2 + 1.4854P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.001$
 $\Delta\rho_{\text{max}} = 1.10 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -1.02 \text{ e } \text{\AA}^{-3}$

Special details

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 $F^2 > 2\sigma(F^2)$ 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)
Br1	0.91668 (4)	0.11235 (2)	0.50601 (4)	0.09330 (19)	
O1	0.4895 (3)	0.41833 (11)	0.5435 (2)	0.0719 (6)	
O2	0.3062 (2)	0.18500 (11)	0.61555 (16)	0.0577 (5)	
O3	0.3634 (2)	0.13165 (11)	0.79229 (19)	0.0597 (5)	
O4	0.4398 (2)	0.31539 (12)	0.38649 (17)	0.0657 (6)	
H4	0.4496	0.3478	0.4434	0.099*	
N1	0.6847 (2)	0.28211 (11)	0.87199 (18)	0.0436 (5)	
H1	0.7348	0.2820	0.9422	0.052*	
C1	0.6826 (2)	0.34138 (12)	0.8040 (2)	0.0392 (5)	
C2	0.7744 (3)	0.40011 (14)	0.8550 (2)	0.0487 (6)	
H2A	0.8640	0.3812	0.8910	0.058*	
H2B	0.7312	0.4228	0.9193	0.058*	
C3	0.8001 (3)	0.45464 (14)	0.7599 (3)	0.0507 (6)	
C4	0.6576 (3)	0.47395 (14)	0.6893 (3)	0.0584 (7)	
H4A	0.6025	0.4986	0.7434	0.070*	
H4B	0.6726	0.5057	0.6238	0.070*	
C5	0.5754 (3)	0.41168 (14)	0.6362 (3)	0.0494 (6)	
C6	0.5958 (2)	0.34543 (12)	0.6974 (2)	0.0385 (5)	
C7	0.5182 (2)	0.28168 (12)	0.64397 (19)	0.0372 (5)	
H7A	0.4239	0.2969	0.6095	0.045*	
C8	0.5009 (2)	0.22835 (12)	0.74306 (19)	0.0350 (5)	
C9	0.5861 (2)	0.22942 (13)	0.85029 (19)	0.0379 (5)	
C10	0.5868 (3)	0.17973 (17)	0.9557 (2)	0.0576 (7)	

H10A	0.5907	0.1324	0.9270	0.086*	
H10B	0.5028	0.1862	0.9930	0.086*	
H10C	0.6674	0.1889	1.0144	0.086*	
C11	0.8970 (3)	0.42483 (18)	0.6729 (3)	0.0673 (8)	
H11A	0.8543	0.3841	0.6329	0.101*	
H11B	0.9855	0.4121	0.7182	0.101*	
H11C	0.9120	0.4596	0.6132	0.101*	
C12	0.8689 (4)	0.51939 (17)	0.8232 (4)	0.0734 (9)	
H12A	0.8085	0.5383	0.8779	0.110*	
H12B	0.8840	0.5540	0.7633	0.110*	
H12C	0.9574	0.5065	0.8684	0.110*	
C13	0.3830 (2)	0.18055 (13)	0.7107 (2)	0.0407 (5)	
C14	0.2502 (7)	0.0825 (4)	0.7444 (6)	0.0617 (18)	0.633 (10)
H14A	0.2704	0.0612	0.6690	0.074*	0.633 (10)
H14B	0.1606	0.1066	0.7295	0.074*	0.633 (10)
C15	0.2477 (6)	0.0278 (4)	0.8428 (7)	0.080 (2)	0.633 (10)
H15A	0.1797	-0.0076	0.8157	0.120*	0.633 (10)
H15B	0.2229	0.0495	0.9154	0.120*	0.633 (10)
H15C	0.3389	0.0067	0.8599	0.120*	0.633 (10)
C14'	0.2263 (8)	0.0963 (5)	0.7967 (10)	0.051 (3)	0.367 (10)
H14C	0.1501	0.1219	0.7502	0.061*	0.367 (10)
H14D	0.2062	0.0910	0.8799	0.061*	0.367 (10)
C15'	0.2497 (11)	0.0255 (6)	0.7387 (15)	0.094 (5)	0.367 (10)
H15D	0.1635	-0.0007	0.7296	0.141*	0.367 (10)
H15E	0.3205	-0.0001	0.7899	0.141*	0.367 (10)
H15F	0.2798	0.0326	0.6602	0.141*	0.367 (10)
C16	0.5888 (3)	0.25025 (13)	0.5398 (2)	0.0407 (5)	
C17	0.6996 (3)	0.20358 (14)	0.5642 (2)	0.0459 (6)	
H17A	0.7331	0.1922	0.6444	0.055*	
C18	0.7603 (3)	0.17389 (15)	0.4702 (3)	0.0561 (7)	
C19	0.7115 (4)	0.18797 (18)	0.3507 (3)	0.0662 (9)	
H19A	0.7507	0.1661	0.2881	0.079*	
C20	0.6038 (4)	0.23495 (18)	0.3256 (2)	0.0643 (9)	
H20A	0.5706	0.2451	0.2448	0.077*	
C21	0.5431 (3)	0.26783 (15)	0.4182 (2)	0.0499 (6)	

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Br1	0.0748 (3)	0.1056 (3)	0.1086 (4)	0.0222 (2)	0.0469 (2)	0.0037 (2)
O1	0.0849 (15)	0.0519 (12)	0.0703 (14)	0.0074 (11)	-0.0226 (12)	0.0177 (10)
O2	0.0550 (11)	0.0705 (13)	0.0419 (10)	-0.0123 (9)	-0.0152 (8)	0.0044 (9)
O3	0.0498 (11)	0.0594 (11)	0.0645 (12)	-0.0181 (9)	-0.0140 (9)	0.0208 (10)
O4	0.0744 (14)	0.0777 (14)	0.0404 (10)	-0.0011 (11)	-0.0102 (9)	0.0150 (10)
N1	0.0421 (10)	0.0523 (12)	0.0332 (9)	-0.0084 (9)	-0.0066 (8)	0.0074 (9)
C1	0.0378 (11)	0.0429 (12)	0.0373 (11)	-0.0015 (9)	0.0062 (9)	0.0009 (10)
C2	0.0483 (14)	0.0512 (15)	0.0459 (14)	-0.0081 (11)	0.0028 (11)	-0.0035 (11)
C3	0.0509 (14)	0.0430 (14)	0.0595 (16)	-0.0037 (11)	0.0123 (12)	-0.0006 (12)
C4	0.0626 (17)	0.0386 (14)	0.0738 (19)	0.0055 (12)	0.0078 (14)	0.0038 (13)
C5	0.0514 (15)	0.0432 (14)	0.0526 (15)	0.0085 (11)	0.0036 (12)	0.0053 (11)

C6	0.0399 (12)	0.0405 (12)	0.0349 (11)	0.0032 (9)	0.0040 (9)	0.0039 (9)
C7	0.0373 (11)	0.0447 (13)	0.0280 (10)	0.0021 (9)	-0.0024 (8)	0.0054 (9)
C8	0.0337 (10)	0.0426 (12)	0.0284 (10)	0.0016 (9)	0.0032 (8)	0.0042 (9)
C9	0.0373 (11)	0.0463 (13)	0.0298 (10)	-0.0021 (10)	0.0031 (8)	0.0047 (9)
C10	0.0631 (17)	0.0708 (18)	0.0352 (12)	-0.0208 (14)	-0.0084 (11)	0.0165 (12)
C11	0.0628 (18)	0.070 (2)	0.074 (2)	-0.0017 (15)	0.0281 (16)	0.0055 (16)
C12	0.075 (2)	0.0544 (18)	0.092 (2)	-0.0175 (16)	0.0133 (18)	-0.0061 (17)
C13	0.0382 (11)	0.0438 (13)	0.0388 (12)	0.0019 (10)	0.0002 (9)	0.0021 (10)
C14	0.061 (3)	0.065 (4)	0.055 (4)	-0.022 (3)	-0.004 (3)	0.005 (3)
C15	0.065 (3)	0.084 (4)	0.093 (5)	-0.018 (3)	0.012 (3)	0.028 (4)
C14'	0.047 (4)	0.062 (6)	0.044 (6)	-0.012 (4)	0.003 (4)	0.006 (4)
C15'	0.053 (5)	0.064 (7)	0.169 (16)	-0.009 (5)	0.025 (7)	-0.002 (8)
C16	0.0462 (13)	0.0448 (13)	0.0311 (11)	-0.0106 (10)	0.0046 (9)	0.0024 (10)
C17	0.0461 (13)	0.0529 (14)	0.0398 (12)	-0.0080 (11)	0.0102 (10)	0.0020 (11)
C18	0.0559 (15)	0.0575 (16)	0.0594 (17)	-0.0096 (13)	0.0248 (13)	-0.0029 (13)
C19	0.081 (2)	0.073 (2)	0.0509 (16)	-0.0240 (18)	0.0309 (15)	-0.0138 (15)
C20	0.081 (2)	0.083 (2)	0.0298 (12)	-0.0262 (18)	0.0102 (13)	-0.0010 (13)
C21	0.0578 (15)	0.0581 (16)	0.0328 (12)	-0.0163 (13)	0.0022 (10)	0.0049 (11)

Geometric parameters (Å, °)

Br1—C18	1.907 (3)	C10—H10A	0.9600
O1—C5	1.238 (3)	C10—H10B	0.9600
O2—C13	1.208 (3)	C10—H10C	0.9600
O3—C13	1.329 (3)	C11—H11A	0.9600
O3—C14	1.481 (5)	C11—H11B	0.9600
O3—C14'	1.485 (7)	C11—H11C	0.9600
O4—C21	1.357 (4)	C12—H12A	0.9600
O4—H4	0.8798	C12—H12B	0.9600
N1—C1	1.357 (3)	C12—H12C	0.9600
N1—C9	1.381 (3)	C14—C15	1.512 (4)
N1—H1	0.8600	C14—H14A	0.9700
C1—C6	1.356 (3)	C14—H14B	0.9700
C1—C2	1.491 (3)	C15—H15A	0.9600
C2—C3	1.523 (4)	C15—H15B	0.9600
C2—H2A	0.9700	C15—H15C	0.9600
C2—H2B	0.9700	C14'—C15'	1.526 (5)
C3—C12	1.528 (4)	C14'—H14C	0.9700
C3—C4	1.530 (4)	C14'—H14D	0.9700
C3—C11	1.531 (4)	C15'—H15D	0.9600
C4—C5	1.503 (4)	C15'—H15E	0.9600
C4—H4A	0.9700	C15'—H15F	0.9600
C4—H4B	0.9700	C16—C17	1.386 (4)
C5—C6	1.436 (3)	C16—C21	1.402 (3)
C6—C7	1.507 (3)	C17—C18	1.378 (4)
C7—C8	1.521 (3)	C17—H17A	0.9300
C7—C16	1.532 (3)	C18—C19	1.371 (4)
C7—H7A	0.9800	C19—C20	1.369 (5)
C8—C9	1.352 (3)	C19—H19A	0.9300
C8—C13	1.461 (3)	C20—C21	1.392 (4)

C9—C10	1.503 (3)	C20—H20A	0.9300
C13—O3—C14	111.3 (3)	C3—C11—H11A	109.5
C13—O3—C14'	123.0 (5)	C3—C11—H11B	109.5
C21—O4—H4	106.1	H11A—C11—H11B	109.5
C1—N1—C9	123.33 (19)	C3—C11—H11C	109.5
C1—N1—H1	118.0	H11A—C11—H11C	109.5
C9—N1—H1	116.5	H11B—C11—H11C	109.5
C6—C1—N1	119.6 (2)	C3—C12—H12A	109.5
C6—C1—C2	123.6 (2)	C3—C12—H12B	109.5
N1—C1—C2	116.8 (2)	H12A—C12—H12B	109.5
C1—C2—C3	113.1 (2)	C3—C12—H12C	109.5
C1—C2—H2A	109.0	H12A—C12—H12C	109.5
C3—C2—H2A	109.0	H12B—C12—H12C	109.5
C1—C2—H2B	109.0	O2—C13—O3	121.2 (2)
C3—C2—H2B	109.0	O2—C13—C8	122.4 (2)
H2A—C2—H2B	107.8	O3—C13—C8	116.37 (19)
C2—C3—C12	109.5 (2)	O3—C14—C15	105.0 (4)
C2—C3—C4	107.7 (2)	O3—C14—H14A	110.8
C12—C3—C4	110.2 (2)	C15—C14—H14A	110.8
C2—C3—C11	110.2 (2)	O3—C14—H14B	110.8
C12—C3—C11	109.1 (3)	C15—C14—H14B	110.8
C4—C3—C11	110.1 (3)	H14A—C14—H14B	108.8
C5—C4—C3	113.5 (2)	O3—C14'—C15'	102.1 (6)
C5—C4—H4A	108.9	O3—C14'—H14C	111.4
C3—C4—H4A	108.9	C15'—C14'—H14C	111.4
C5—C4—H4B	108.9	O3—C14'—H14D	111.4
C3—C4—H4B	108.9	C15'—C14'—H14D	111.4
H4A—C4—H4B	107.7	H14C—C14'—H14D	109.2
O1—C5—C6	121.2 (3)	C14'—C15'—H15D	109.5
O1—C5—C4	120.2 (2)	C14'—C15'—H15E	109.5
C6—C5—C4	118.6 (2)	H15D—C15'—H15E	109.5
C1—C6—C5	119.5 (2)	C14'—C15'—H15F	109.5
C1—C6—C7	120.9 (2)	H15D—C15'—H15F	109.5
C5—C6—C7	119.7 (2)	H15E—C15'—H15F	109.5
C6—C7—C8	110.56 (18)	C17—C16—C21	118.5 (2)
C6—C7—C16	111.54 (19)	C17—C16—C7	120.5 (2)
C8—C7—C16	112.36 (19)	C21—C16—C7	121.0 (2)
C6—C7—H7A	107.4	C18—C17—C16	120.4 (2)
C8—C7—H7A	107.4	C18—C17—H17A	119.8
C16—C7—H7A	107.4	C16—C17—H17A	119.8
C9—C8—C13	125.9 (2)	C19—C18—C17	121.5 (3)
C9—C8—C7	120.9 (2)	C19—C18—Br1	118.9 (2)
C13—C8—C7	113.17 (18)	C17—C18—Br1	119.6 (2)
C8—C9—N1	119.2 (2)	C20—C19—C18	118.7 (3)
C8—C9—C10	127.9 (2)	C20—C19—H19A	120.7
N1—C9—C10	112.93 (19)	C18—C19—H19A	120.7
C9—C10—H10A	109.5	C19—C20—C21	121.5 (3)
C9—C10—H10B	109.5	C19—C20—H20A	119.3

H10A—C10—H10B	109.5	C21—C20—H20A	119.3
C9—C10—H10C	109.5	O4—C21—C20	118.2 (2)
H10A—C10—H10C	109.5	O4—C21—C16	122.4 (2)
H10B—C10—H10C	109.5	C20—C21—C16	119.4 (3)
C9—N1—C1—C6	-11.6 (4)	C1—N1—C9—C8	14.5 (4)
C9—N1—C1—C2	166.2 (2)	C1—N1—C9—C10	-164.2 (2)
C6—C1—C2—C3	-19.6 (4)	C14—O3—C13—O2	6.0 (5)
N1—C1—C2—C3	162.7 (2)	C14'—O3—C13—O2	-21.9 (6)
C1—C2—C3—C12	168.6 (2)	C14—O3—C13—C8	-174.5 (4)
C1—C2—C3—C4	48.8 (3)	C14'—O3—C13—C8	157.6 (5)
C1—C2—C3—C11	-71.4 (3)	C9—C8—C13—O2	175.8 (2)
C2—C3—C4—C5	-53.9 (3)	C7—C8—C13—O2	-1.8 (3)
C12—C3—C4—C5	-173.3 (3)	C9—C8—C13—O3	-3.7 (4)
C11—C3—C4—C5	66.4 (3)	C7—C8—C13—O3	178.8 (2)
C3—C4—C5—O1	-152.6 (3)	C13—O3—C14—C15	176.1 (5)
C3—C4—C5—C6	29.2 (4)	C14'—O3—C14—C15	-62.6 (10)
N1—C1—C6—C5	169.3 (2)	C13—O3—C14'—C15'	102.7 (10)
C2—C1—C6—C5	-8.3 (4)	C14—O3—C14'—C15'	31.1 (9)
N1—C1—C6—C7	-9.4 (3)	C6—C7—C16—C17	84.8 (3)
C2—C1—C6—C7	173.0 (2)	C8—C7—C16—C17	-40.0 (3)
O1—C5—C6—C1	-174.8 (3)	C6—C7—C16—C21	-95.2 (3)
C4—C5—C6—C1	3.3 (4)	C8—C7—C16—C21	140.0 (2)
O1—C5—C6—C7	3.9 (4)	C21—C16—C17—C18	-1.8 (4)
C4—C5—C6—C7	-178.0 (2)	C7—C16—C17—C18	178.2 (2)
C1—C6—C7—C8	24.1 (3)	C16—C17—C18—C19	-1.6 (4)
C5—C6—C7—C8	-154.6 (2)	C16—C17—C18—Br1	177.85 (19)
C1—C6—C7—C16	-101.7 (2)	C17—C18—C19—C20	2.8 (4)
C5—C6—C7—C16	79.6 (3)	Br1—C18—C19—C20	-176.7 (2)
C6—C7—C8—C9	-21.1 (3)	C18—C19—C20—C21	-0.5 (5)
C16—C7—C8—C9	104.2 (2)	C19—C20—C21—O4	178.0 (3)
C6—C7—C8—C13	156.56 (19)	C19—C20—C21—C16	-2.8 (4)
C16—C7—C8—C13	-78.1 (2)	C17—C16—C21—O4	-177.0 (2)
C13—C8—C9—N1	-173.8 (2)	C7—C16—C21—O4	3.1 (4)
C7—C8—C9—N1	3.6 (3)	C17—C16—C21—C20	3.9 (4)
C13—C8—C9—C10	4.8 (4)	C7—C16—C21—C20	-176.0 (2)
C7—C8—C9—C10	-177.8 (3)		

Hydrogen-bond geometry (Å, °)

<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>
O4—H4...O1	0.88	1.75	2.625 (3)	171
N1—H1...O2 ⁱ	0.86	2.05	2.866 (3)	158

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