



Received 12 September 2015
Accepted 16 September 2015

Edited by A. J. Lough, University of Toronto,
Canada

Keywords: crystal structure; bromanilic acid;
morpholine; hydrogen-bonding; proton disorder

CCDC references: 1424714; 1424713;
1424712

Supporting information: this article has
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Crystal structures of morpholinium hydrogen bromanilate at 130, 145 and 180 K

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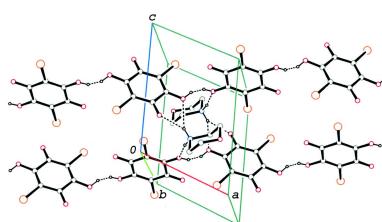
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Crystal structures of the title compound (systematic name: morpholin-4-i um 2,5-dibromo-4-hydroxy-3,6-dioxocyclohexa-1,4-dien-1-olate), $C_4H_{10}NO^+ \cdot C_6HBr_2O_4^-$, were determined at three temperatures, *viz.* 130, 145 and 180 K. The asymmetric unit comprises one morpholinium cation and two halves of crystallographically independent bromanilate monoanions, which are located on inversion centres. The conformations of the two independent bromanilate anions are different from each other with respect to the O—H orientation. In the crystal, the two different anions are linked alternately into a chain along [211] through a short O—H \cdots O hydrogen bond, in which the H atom is disordered over two positions. The refined site-occupancy ratios, which are almost constant in the temperature range studied, are 0.49 (3):0.51 (3), 0.52 (3):0.48 (3) and 0.50 (3):0.50 (3), respectively, at 130, 145 and 180 K, and no significant difference in the molecular geometry and the molecular packing is observed at the three temperatures. The morpholinium cation links adjacent chains of anions *via* N—H \cdots O hydrogen bonds, forming a sheet structure parallel to (111).

1. Chemical context

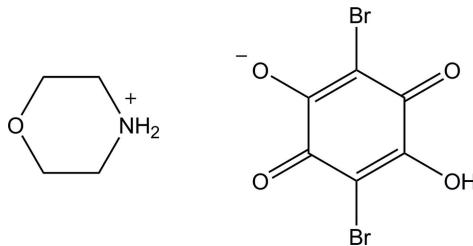
Anilic acid (2,5-dihydroxy-1,4-benzoquinone) derivatives, such as chloranilic acid (2,5-dichloro-3,6-dihydroxy-1,4-benzoquinone) and bromanilic acid (2,5-dibromo-3,6-dihydroxy-1,4-benzoquinone), appear particularly attractive as a versatile template for generating hydrogen-bonded self-assemblies with various organic bases (Zaman *et al.*, 2001; Molčanov & Kojić-Prodić, 2010; Gotoh & Ishida, 2011; Thomas *et al.*, 2013) and also as a model compound for investigating proton dynamics in hydrogen-bond systems (Ikeda *et al.*, 2005; Seliger *et al.*, 2009). Furthermore, salts and co-crystals of anilic acids with organic bases have attracted much interest with respect to organic ferroelectrics (Horiuchi *et al.*, 2008, 2009, 2013).

In our previous study, we reported the crystal structure of morpholinium hydrogen chloranilate, $C_4H_{10}NO^+ \cdot C_6HCl_2O_4^-$, in which a short O—H \cdots O hydrogen bond is formed between the chloranilate ions and the H atom in the hydrogen bond is disordered over two sites (Ishida & Kashino, 1999). The measurements of ^{35}Cl NQR (nuclear quadrupole resonance) for the compound in the temperature range 4–300 K showed an anomalous temperature dependence of the NQR frequencies, which cannot be explained by the conventional Bayer-type lattice motion: one of the two frequencies exhibits an anomalous increase with increasing temperature from 4.2 K while the other frequency shows a rather fast decrease with temperature. The anomalous behavior was ascribed to a drastic temperature variation of the disordered O—H \cdots O



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hydrogen bond, as revealed by multi-temperature X-ray diffraction (Tobu *et al.*, 2012). In the present study, we have undertaken the structural determination of morpholinium hydrogen bromanilate, $C_4H_{10}NO^+ \cdot C_6HBr_2O_4^-$, to extend the study of hydrogen-bonding in the amine-halohydroxybenzoquinone system.



2. Structural commentary

The title compound is isomorphous with morpholinium hydrogen chloranilate in the space group $P\bar{1}$ (Ishida & Kashino, 1999; Tobu *et al.*, 2012) and has a quite similar molecular packing to the chloranilate. The asymmetric unit of the title compound comprises one morpholinium cation and two halves of crystallographically independent bromanilate monoanions, which are each located on an inversion centre (Fig. 1). The conformations of two bromanilate anions are different from each other with respect to the O—H orientation as shown schematically in Fig. 2.

In morpholinium hydrogen chloranilate, the bond distances of C3—O2 and C6—O4, which are involved in the disordered O—H \cdots O hydrogen bond, showed slight but systematic decrease and increase, respectively, with temperature [C3—O2: from 1.2994 (10) Å at 114 K to 1.2951 (10) Å at 180 K;

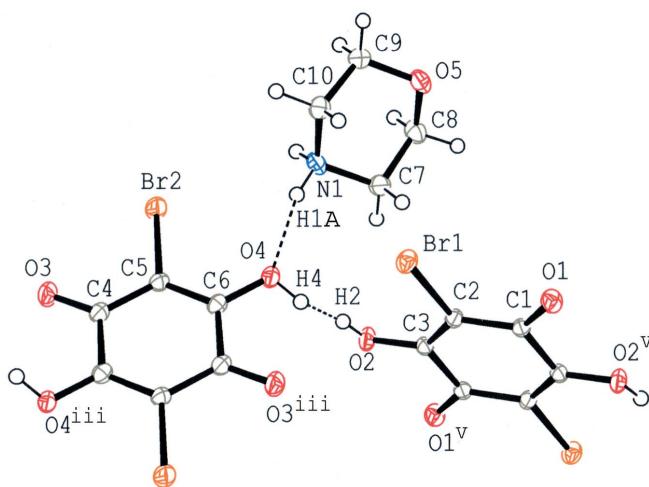


Figure 1

A view of the molecular structure of the title compound at 180 K, showing the atom-numbering scheme. Displacement ellipsoids of non-H atoms are drawn at the 50% probability level and H atoms are drawn as circles of arbitrary size. The site-occupancy factors of the disordered H atom (H2 and H4) are approximately equal. The N—H \cdots O and O—H \cdots O hydrogen bonds are indicated by dashed lines. [Symmetry codes: (iii) $-x, -y + 1, -z$; (v) $-x + 2, -y + 2, -z + 1$.]

Table 1
Hydrogen-bond geometry (Å, °) at 130 K.

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N1—H1A \cdots O4	0.88 (3)	2.03 (3)	2.886 (2)	166 (2)
N1—H1B \cdots O1 ⁱ	0.86 (3)	2.16 (3)	2.938 (2)	150 (2)
N1—H1B \cdots O2 ⁱⁱ	0.86 (3)	2.27 (3)	2.955 (2)	137 (2)
O2—H2 \cdots O4	0.81 (3)	1.77 (3)	2.5160 (16)	152 (4)
O2—H2 \cdots O3 ⁱⁱⁱ	0.81 (3)	2.57 (3)	3.0613 (17)	120 (3)
O4—H4 \cdots O2	0.82 (3)	1.82 (4)	2.5160 (16)	143 (4)
C7—H7A \cdots O4 ⁱⁱ	0.99	2.53	3.391 (2)	145
C10—H10B \cdots Br2 ^{iv}	0.99	2.90	3.8892 (17)	175

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

Table 2
Hydrogen-bond geometry (Å, °) at 145 K.

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N1—H1A \cdots O4	0.86 (3)	2.04 (3)	2.888 (2)	166 (2)
N1—H1B \cdots O1 ⁱ	0.85 (3)	2.17 (3)	2.938 (2)	151 (2)
N1—H1B \cdots O2 ⁱⁱ	0.85 (3)	2.29 (3)	2.959 (2)	136 (2)
O2—H2 \cdots O4	0.82 (3)	1.77 (3)	2.5174 (16)	153 (4)
O2—H2 \cdots O3 ⁱⁱⁱ	0.82 (3)	2.58 (3)	3.0628 (17)	120 (3)
O4—H4 \cdots O2	0.82 (3)	1.79 (4)	2.5174 (16)	147 (4)
C7—H7A \cdots O4 ⁱⁱ	0.99	2.54	3.394 (2)	145
C10—H10B \cdots Br2 ^{iv}	0.99	2.90	3.8905 (17)	175

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

Table 3
Hydrogen-bond geometry (Å, °) at 180 K.

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N1—H1A \cdots O4	0.89 (3)	2.02 (3)	2.890 (2)	167 (2)
N1—H1B \cdots O1 ⁱ	0.86 (3)	2.16 (3)	2.938 (2)	150 (2)
N1—H1B \cdots O2 ⁱⁱ	0.86 (3)	2.28 (3)	2.964 (2)	136 (2)
O2—H2 \cdots O4	0.82 (3)	1.79 (4)	2.5224 (16)	148 (5)
O2—H2 \cdots O3 ⁱⁱⁱ	0.82 (3)	2.55 (4)	3.0678 (18)	122 (4)
O4—H4 \cdots O2	0.82 (3)	1.80 (4)	2.5224 (16)	147 (4)
C7—H7A \cdots O4 ⁱⁱ	0.99	2.55	3.402 (2)	145
C10—H10B \cdots Br2 ^{iv}	0.99	2.91	3.8946 (17)	174

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

C6—O4: from 1.290 (10) Å at 114 K to 1.2946 (10) Å at 180 K], which corresponds to population changes of the two disordered proton sites in the hydrogen bond (Tobu *et al.*, 2012). In the present compound, however, the C3—O2 and C6—O4 bond lengths are almost constant [C3—O2: 1.2953 (17),

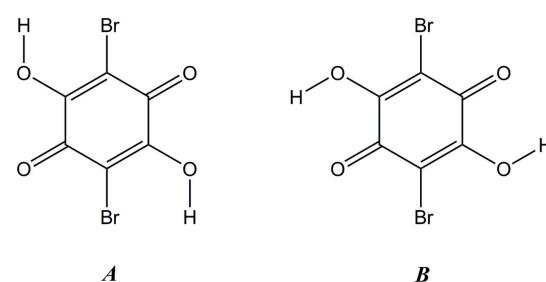
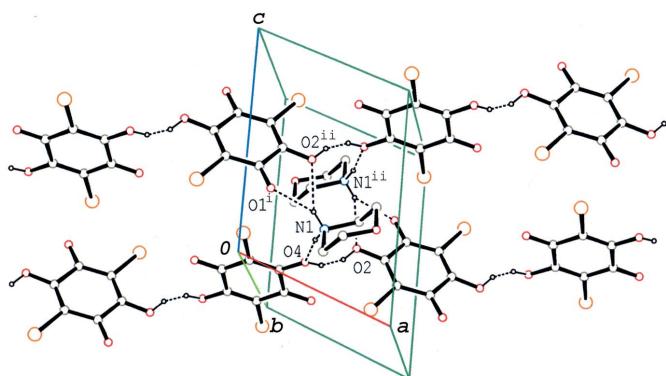


Figure 2

Two conformations (A and B forms) of bromanilic acid with respect to the O—H orientation.

**Figure 3**

A partial packing diagram of the title compound at 180 K, showing the hydrogen-bonded aggregate of morpholinium and hydrogen bromanilate ions. The N—H···O and O—H···O hydrogen bonds are indicated by dashed lines. [Symmetry codes: (i) $x - 1, y - 1, z$; (ii) $-x + 1, -y + 1, -z + 1$.]

1.2937 (17) and 1.2931 (17) Å at 130, 145 and 180 K; C6—O4: 1.3002 (18), 1.2997 (18) and 1.2997 (18) Å at 130, 145 and 180 K] and no significant difference in the molecular geometry is observed at the three temperatures.

3. Supramolecular features

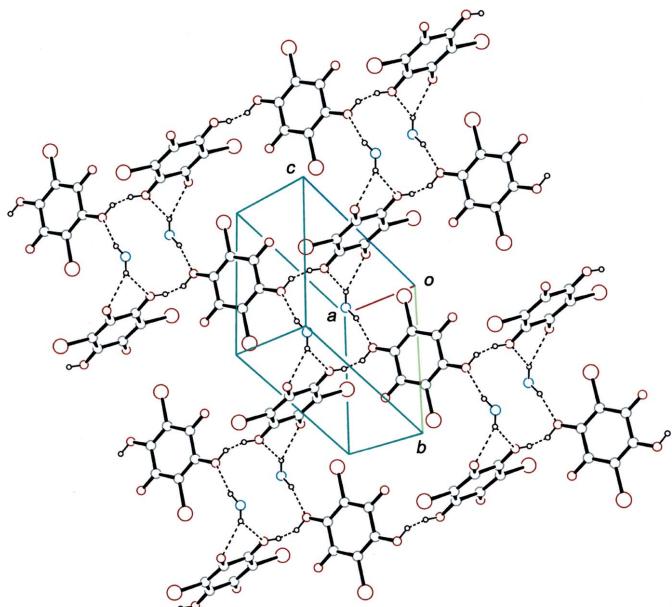
In the crystal, the two independent bromanilate anions with different conformations are linked alternately by short O—H···O hydrogen bonds (Tables 1, 2 and 3), forming a chain along [211] (Fig. 3). The adjacent independent anions are almost perpendicular to each other, with dihedral angles of 86.57 (7)° (130 K), 86.65 (7)° (145 K) and 86.81 (7)° (180 K) between the benzoquinone rings. The morpholinium cation connects the anion chains through N—H···O hydrogen bonds and a weak C—H···O hydrogen bond into a sheet parallel to (111) (Fig. 4). Between the chains, short Br···O and Br···C contacts [Br₂···O1ⁱ: 3.1698 (13) Å (130 K), 3.1725 (13) Å (145 K) and 3.1763 (13) Å (180 K); Br₂···C1ⁱ: 3.2673 (15) Å (130 K), 3.2716 (15) Å (145 K) and 3.2808 (15) Å (180 K); symmetry code: (i) $x - 1, y - 1, z$] are observed. A weak C···Br interaction is also observed between the sheets.

4. Database survey

Although a search of the Cambridge Structural Database (Version 5.36, last update February 2015; Groom & Allen, 2014) for organic salts and co-crystals with bromanilic acid gave 31 hits, no crystal structure including the A form (Fig. 2) was found.

5. Synthesis and crystallization

Single crystals of the title compound suitable for X-ray diffraction were prepared by slow evaporation from an acetonitrile solution (200 ml) of bromanilic acid (200 mg) with morpholine (60 mg) at room temperature.

**Figure 4**

A packing diagram of the title compound at 180 K, showing the sheet structure formed through N—H···O and O—H···O hydrogen bonds (dashed lines). For the morpholinium cations, only NH₂ groups are shown for clarity.

6. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 4. C-bound H atoms of the morpholinium cation were positioned geometrically with C—H = 0.99 Å and were refined as riding with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$. The N-bound H atom was located in a difference Fourier map and refined freely [refined N—H = 0.85 (3)–0.89 (3) Å]. Two disordered positions of the H atom in the O—H···O hydrogen bond were located in a difference Fourier map. Since site occupancy factors and isotropic displacement parameters are correlated and bonding effects also make the site-occupancy factors less reliable, the positional parameters and the occupancies of the H atom were refined, with $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{O})$, and with distance restraints of O—H = 0.84 (2) Å.

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Table 4
Experimental details.

	130 K	145 K	180 K
Crystal data			
Chemical formula	$C_4H_{10}NO^+ \cdot C_6HBr_2O_4^-$	$C_4H_{10}NO^+ \cdot C_6HBr_2O_4^-$	$C_4H_{10}NO^+ \cdot C_6HBr_2O_4^-$
M_r	385.01	385.01	385.01
Crystal system, space group	Triclinic, $P\bar{1}$	Triclinic, $P\bar{1}$	Triclinic, $P\bar{1}$
a, b, c (Å)	8.62046 (19), 9.2129 (2), 9.4257 (2)	8.62293 (18), 9.21849 (19), 9.4354 (2)	8.62824 (17), 9.23087 (18), 9.46007 (19)
α, β, γ (°)	93.5208 (7), 112.9139 (7), 115.9757 (7)	93.5239 (7), 112.9190 (7), 115.9777 (7)	93.5321 (7), 112.9738 (7), 115.9508 (7)
V (Å ³)	595.05 (3)	596.13 (3)	598.67 (3)
Z	2	2	2
Radiation type	Mo $K\alpha$	Mo $K\alpha$	Mo $K\alpha$
μ (mm ⁻¹)	6.84	6.83	6.80
Crystal size (mm)	0.40 × 0.34 × 0.18	0.40 × 0.34 × 0.18	0.40 × 0.34 × 0.18
Data collection			
Diffractometer	Rigaku R-AXIS RAPIDII	Rigaku R-AXIS RAPIDII	Rigaku R-AXIS RAPIDII
Absorption correction	Numerical (<i>NUMABS</i> ; Higashi, 1999)	Numerical (<i>NUMABS</i> ; Higashi, 1999)	Numerical (<i>NUMABS</i> ; Higashi, 1999)
T_{\min}, T_{\max}	0.096, 0.292	0.098, 0.292	0.098, 0.294
No. of measured, independent and observed [$I > 2\sigma(I)$] reflections	18162, 3468, 3183	18176, 3473, 3181	18199, 3487, 3188
R_{int}	0.026	0.028	0.026
(sin θ/λ) _{max} (Å ⁻¹)	0.704	0.704	0.703
Refinement			
$R[F^2 > 2\sigma(F^2)], wR(F^2), S$	0.017, 0.046, 1.14	0.018, 0.046, 1.10	0.019, 0.048, 1.09
No. of reflections	3468	3473	3487
No. of parameters	178	178	178
No. of restraints	2	2	2
H-atom treatment	H atoms treated by a mixture of independent and constrained refinement	H atoms treated by a mixture of independent and constrained refinement	H atoms treated by a mixture of independent and constrained refinement
$\Delta\rho_{\max}, \Delta\rho_{\min}$ (e Å ⁻³)	0.50, -0.37	0.48, -0.44	0.59, -0.45

Computer programs: *RAPID-AUTO* (Rigaku, 2006), *SIR92* (Altomare *et al.*, 1994), *SHELXL2014/7* (Sheldrick, 2015), *ORTEP-3 for Windows* (Farrugia, 2012), *CrystalStructure* (Rigaku, 2010) and *PLATON* (Spek, 2009).

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supporting information

Acta Cryst. (2015). E71, 1226-1229 [doi:10.1107/S2056989015017272]

Crystal structures of morpholinium hydrogen bromanilate at 130, 145 and 180 K

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Computing details

For all compounds, data collection: *RAPID-AUTO* (Rigaku, 2006); cell refinement: *RAPID-AUTO* (Rigaku, 2006); data reduction: *RAPID-AUTO* (Rigaku, 2006); program(s) used to solve structure: *SIR92* (Altomare *et al.*, 1994). Program(s) used to refine structure: *SHELXL2014* (Sheldrick, 2015) for (1); *SHELXL2014/7* (Sheldrick, 2015) for (2), (3). For all compounds, molecular graphics: *ORTEP-3 for Windows* (Farrugia, 2012); software used to prepare material for publication: *CrystalStructure* (Rigaku, 2010) and *PLATON* (Spek, 2009).

(1) Morpholin-4-ium 2,5-dibromo-4-hydroxy-3,6-dioxocyclohexa-1,4-dien-1-olate

Crystal data

$\text{C}_4\text{H}_{10}\text{NO}^+\cdot\text{C}_6\text{HBr}_2\text{O}_4^-$	$Z = 2$
$M_r = 385.01$	$F(000) = 376.00$
Triclinic, $P\bar{1}$	$D_x = 2.149 \text{ Mg m}^{-3}$
Hall symbol: -P 1	Mo $K\alpha$ radiation, $\lambda = 0.71075 \text{ \AA}$
$a = 8.62046 (19) \text{ \AA}$	Cell parameters from 15996 reflections
$b = 9.2129 (2) \text{ \AA}$	$\theta = 3.0\text{--}30.1^\circ$
$c = 9.4257 (2) \text{ \AA}$	$\mu = 6.84 \text{ mm}^{-1}$
$\alpha = 93.5208 (7)^\circ$	$T = 130 \text{ K}$
$\beta = 112.9139 (7)^\circ$	Block, brown
$\gamma = 115.9757 (7)^\circ$	$0.40 \times 0.34 \times 0.18 \text{ mm}$
$V = 595.05 (3) \text{ \AA}^3$	

Data collection

Rigaku R-AXIS RAPIDII	3468 independent reflections
diffractometer	3183 reflections with $I > 2\sigma(I)$
Detector resolution: 10.000 pixels mm^{-1}	$R_{\text{int}} = 0.026$
ω scans	$\theta_{\text{max}} = 30.0^\circ, \theta_{\text{min}} = 3.0^\circ$
Absorption correction: numerical (NUMABS; Higashi, 1999)	$h = -12 \rightarrow 12$
$T_{\text{min}} = 0.096, T_{\text{max}} = 0.292$	$k = -12 \rightarrow 12$
18162 measured reflections	$l = -13 \rightarrow 13$

Refinement

Refinement on F^2	178 parameters
Least-squares matrix: full	2 restraints
$R[F^2 > 2\sigma(F^2)] = 0.017$	Hydrogen site location: mixed
$wR(F^2) = 0.046$	H atoms treated by a mixture of independent and constrained refinement
$S = 1.14$	
3468 reflections	

$$w = 1/[\sigma^2(F_o^2) + (0.0195P)^2 + 0.3723P]$$

where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.002$

$$\Delta\rho_{\max} = 0.50 \text{ e } \text{\AA}^{-3}$$

$$\Delta\rho_{\min} = -0.37 \text{ e } \text{\AA}^{-3}$$

Special details

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. Reflections were merged by *SHELXL* according to the crystal class for the calculation of statistics and refinement.

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	0.75783 (2)	0.69637 (2)	0.16055 (2)	0.01583 (4)	
Br2	-0.01546 (2)	0.16774 (2)	0.12333 (2)	0.01833 (4)	
O1	1.18246 (16)	0.98222 (14)	0.33171 (13)	0.0166 (2)	
O2	0.62026 (15)	0.76781 (13)	0.41041 (13)	0.0157 (2)	
H2	0.548 (5)	0.699 (4)	0.323 (3)	0.023*	0.49 (3)
O3	-0.34652 (16)	0.22816 (14)	-0.08651 (15)	0.0209 (2)	
O4	0.33771 (16)	0.50578 (13)	0.19213 (13)	0.0164 (2)	
H4	0.427 (5)	0.603 (3)	0.226 (5)	0.025*	0.51 (3)
O5	0.82461 (16)	0.26388 (15)	0.42254 (14)	0.0201 (2)	
N1	0.48737 (19)	0.29440 (17)	0.32853 (17)	0.0170 (2)	
C1	1.0924 (2)	0.98617 (17)	0.40452 (17)	0.0123 (2)	
C2	0.8902 (2)	0.86739 (17)	0.35299 (17)	0.0124 (2)	
C3	0.7974 (2)	0.87220 (17)	0.44093 (17)	0.0126 (2)	
C4	-0.1856 (2)	0.34950 (18)	-0.04567 (18)	0.0146 (3)	
C5	-0.0041 (2)	0.35821 (17)	0.05629 (17)	0.0139 (3)	
C6	0.1730 (2)	0.49677 (18)	0.10262 (17)	0.0145 (3)	
C7	0.6812 (2)	0.42756 (19)	0.45862 (19)	0.0176 (3)	
H7A	0.6639	0.4841	0.5393	0.021*	
H7B	0.7493	0.5130	0.4132	0.021*	
C8	0.8010 (2)	0.3481 (2)	0.53723 (19)	0.0188 (3)	
H8A	0.9304	0.4362	0.6229	0.023*	
H8B	0.7355	0.2669	0.5872	0.023*	
C9	0.6388 (2)	0.1307 (2)	0.3035 (2)	0.0206 (3)	
H9A	0.5750	0.0516	0.3557	0.025*	
H9B	0.6567	0.0681	0.2277	0.025*	
C10	0.5099 (2)	0.1976 (2)	0.21215 (19)	0.0196 (3)	
H10A	0.5690	0.2717	0.1543	0.023*	
H10B	0.3808	0.1032	0.1324	0.023*	
H1A	0.423 (3)	0.342 (3)	0.277 (3)	0.023 (5)*	
H1B	0.422 (3)	0.223 (3)	0.367 (3)	0.027 (6)*	

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Br1	0.01430 (7)	0.01537 (7)	0.01367 (7)	0.00509 (6)	0.00595 (5)	0.00068 (5)
Br2	0.01566 (8)	0.01334 (7)	0.02186 (8)	0.00595 (6)	0.00607 (6)	0.00785 (6)
O1	0.0150 (5)	0.0183 (5)	0.0179 (5)	0.0075 (4)	0.0102 (4)	0.0046 (4)
O2	0.0100 (5)	0.0154 (5)	0.0166 (5)	0.0031 (4)	0.0059 (4)	0.0020 (4)
O3	0.0129 (5)	0.0152 (5)	0.0277 (6)	0.0040 (4)	0.0065 (4)	0.0072 (4)
O4	0.0122 (5)	0.0129 (5)	0.0182 (5)	0.0048 (4)	0.0036 (4)	0.0037 (4)
O5	0.0151 (5)	0.0241 (6)	0.0217 (5)	0.0117 (5)	0.0073 (4)	0.0036 (4)
N1	0.0139 (6)	0.0193 (6)	0.0223 (6)	0.0101 (5)	0.0096 (5)	0.0109 (5)
C1	0.0117 (6)	0.0115 (6)	0.0141 (6)	0.0063 (5)	0.0056 (5)	0.0052 (5)
C2	0.0109 (6)	0.0119 (6)	0.0120 (6)	0.0047 (5)	0.0044 (5)	0.0022 (5)
C3	0.0106 (6)	0.0122 (6)	0.0145 (6)	0.0064 (5)	0.0045 (5)	0.0049 (5)
C4	0.0152 (7)	0.0125 (6)	0.0155 (6)	0.0060 (5)	0.0080 (5)	0.0028 (5)
C5	0.0147 (6)	0.0115 (6)	0.0153 (6)	0.0062 (5)	0.0070 (5)	0.0048 (5)
C6	0.0158 (7)	0.0129 (6)	0.0133 (6)	0.0058 (5)	0.0072 (5)	0.0023 (5)
C7	0.0180 (7)	0.0161 (7)	0.0198 (7)	0.0090 (6)	0.0093 (6)	0.0052 (6)
C8	0.0170 (7)	0.0231 (7)	0.0167 (7)	0.0118 (6)	0.0064 (6)	0.0052 (6)
C9	0.0189 (7)	0.0171 (7)	0.0232 (8)	0.0099 (6)	0.0068 (6)	0.0032 (6)
C10	0.0174 (7)	0.0182 (7)	0.0179 (7)	0.0086 (6)	0.0043 (6)	0.0019 (6)

Geometric parameters (\AA , ^\circ)

Br1—C2	1.8807 (14)	C1—C3 ⁱ	1.5265 (19)
Br2—C5	1.8767 (14)	C2—C3	1.368 (2)
O1—C1	1.2295 (18)	C4—C5	1.445 (2)
O2—C3	1.2953 (17)	C4—C6 ⁱⁱ	1.520 (2)
O2—H2	0.81 (3)	C5—C6	1.362 (2)
O3—C4	1.2229 (18)	C7—C8	1.511 (2)
O4—C6	1.3002 (18)	C7—H7A	0.9900
O4—H4	0.82 (3)	C7—H7B	0.9900
O5—C8	1.4212 (19)	C8—H8A	0.9900
O5—C9	1.428 (2)	C8—H8B	0.9900
N1—C7	1.492 (2)	C9—C10	1.509 (2)
N1—C10	1.494 (2)	C9—H9A	0.9900
N1—H1A	0.88 (3)	C9—H9B	0.9900
N1—H1B	0.86 (3)	C10—H10A	0.9900
C1—C2	1.4407 (19)	C10—H10B	0.9900
C3—O2—H2	118 (3)	C5—C6—C4 ⁱⁱ	120.03 (13)
C6—O4—H4	111 (3)	N1—C7—C8	109.20 (12)
C8—O5—C9	109.71 (12)	N1—C7—H7A	109.8
C7—N1—C10	110.91 (12)	C8—C7—H7A	109.8
C7—N1—H1A	109.1 (14)	N1—C7—H7B	109.8
C10—N1—H1A	108.4 (14)	C8—C7—H7B	109.8
C7—N1—H1B	111.0 (15)	H7A—C7—H7B	108.3
C10—N1—H1B	107.1 (15)	O5—C8—C7	110.61 (13)

H1A—N1—H1B	110 (2)	O5—C8—H8A	109.5
O1—C1—C2	124.43 (13)	C7—C8—H8A	109.5
O1—C1—C3 ⁱ	117.51 (13)	O5—C8—H8B	109.5
C2—C1—C3 ⁱ	118.06 (12)	C7—C8—H8B	109.5
C3—C2—C1	122.59 (13)	H8A—C8—H8B	108.1
C3—C2—Br1	120.39 (11)	O5—C9—C10	111.23 (13)
C1—C2—Br1	116.98 (10)	O5—C9—H9A	109.4
O2—C3—C2	127.82 (13)	C10—C9—H9A	109.4
O2—C3—C1 ⁱ	112.91 (12)	O5—C9—H9B	109.4
C2—C3—C1 ⁱ	119.28 (12)	C10—C9—H9B	109.4
O3—C4—C5	124.23 (14)	H9A—C9—H9B	108.0
O3—C4—C6 ⁱⁱ	118.64 (13)	N1—C10—C9	108.70 (13)
C5—C4—C6 ⁱⁱ	117.13 (12)	N1—C10—H10A	109.9
C6—C5—C4	122.84 (13)	C9—C10—H10A	109.9
C6—C5—Br2	119.19 (11)	N1—C10—H10B	109.9
C4—C5—Br2	117.97 (10)	C9—C10—H10B	109.9
O4—C6—C5	123.65 (14)	H10A—C10—H10B	108.3
O4—C6—C4 ⁱⁱ	116.29 (13)		
O1—C1—C2—C3	177.29 (14)	C6 ⁱⁱ —C4—C5—Br2	-178.24 (10)
C3 ⁱ —C1—C2—C3	-3.3 (2)	C4—C5—C6—O4	-178.83 (14)
O1—C1—C2—Br1	-0.27 (19)	Br2—C5—C6—O4	-0.1 (2)
C3 ⁱ —C1—C2—Br1	179.12 (9)	C4—C5—C6—C4 ⁱⁱ	-0.5 (2)
C1—C2—C3—O2	-176.34 (14)	Br2—C5—C6—C4 ⁱⁱ	178.22 (10)
Br1—C2—C3—O2	1.1 (2)	C10—N1—C7—C8	-54.86 (16)
C1—C2—C3—C1 ⁱ	3.4 (2)	C9—O5—C8—C7	-62.32 (16)
Br1—C2—C3—C1 ⁱ	-179.16 (9)	N1—C7—C8—O5	58.54 (17)
O3—C4—C5—C6	-178.49 (15)	C8—O5—C9—C10	62.50 (17)
C6 ⁱⁱ —C4—C5—C6	0.5 (2)	C7—N1—C10—C9	54.40 (17)
O3—C4—C5—Br2	2.7 (2)	O5—C9—C10—N1	-58.02 (17)

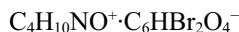
Symmetry codes: (i) $-x+2, -y+2, -z+1$; (ii) $-x, -y+1, -z$.

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

$D\text{—H}\cdots A$	$D\text{—H}$	$H\cdots A$	$D\cdots A$	$D\text{—H}\cdots A$
N1—H1A…O4	0.88 (3)	2.03 (3)	2.886 (2)	166 (2)
N1—H1B…O1 ⁱⁱⁱ	0.86 (3)	2.16 (3)	2.938 (2)	150 (2)
N1—H1B…O2 ^{iv}	0.86 (3)	2.27 (3)	2.955 (2)	137 (2)
O2—H2…O4	0.81 (3)	1.77 (3)	2.5160 (16)	152 (4)
O2—H2…O3 ⁱⁱ	0.81 (3)	2.57 (3)	3.0613 (17)	120 (3)
O4—H4…O2	0.82 (3)	1.82 (4)	2.5160 (16)	143 (4)
C7—H7A…O4 ^{iv}	0.99	2.53	3.391 (2)	145
C10—H10B…Br2 ^v	0.99	2.90	3.8892 (17)	175

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

(2)

Crystal data
 $M_r = 385.01$
Triclinic, $P\bar{1}$

Hall symbol: -P 1

 $a = 8.62293 (18) \text{\AA}$
 $b = 9.21849 (19) \text{\AA}$
 $c = 9.4354 (2) \text{\AA}$
 $\alpha = 93.5239 (7)^\circ$
 $\beta = 112.9190 (7)^\circ$
 $\gamma = 115.9777 (7)^\circ$
 $V = 596.13 (3) \text{\AA}^3$
 $Z = 2$
 $F(000) = 376.00$
 $D_x = 2.145 \text{ Mg m}^{-3}$
Mo $K\alpha$ radiation, $\lambda = 0.71075 \text{\AA}$

Cell parameters from 15916 reflections

 $\theta = 3.0\text{--}30.1^\circ$
 $\mu = 6.83 \text{ mm}^{-1}$
 $T = 145 \text{ K}$

Block, brown

 $0.40 \times 0.34 \times 0.18 \text{ mm}$
*Data collection*Rigaku R-AXIS RAPIDII
diffractometerDetector resolution: 10.000 pixels mm^{-1} ω scansAbsorption correction: numerical
(NUMABS; Higashi, 1999)
 $T_{\min} = 0.098, T_{\max} = 0.292$

18176 measured reflections

3473 independent reflections

3181 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.028$
 $\theta_{\max} = 30.0^\circ, \theta_{\min} = 3.0^\circ$
 $h = -12 \rightarrow 12$
 $k = -12 \rightarrow 12$
 $l = -13 \rightarrow 13$
*Refinement*Refinement on F^2

Least-squares matrix: full

 $R[F^2 > 2\sigma(F^2)] = 0.018$
 $wR(F^2) = 0.046$
 $S = 1.10$

3473 reflections

178 parameters

2 restraints

Hydrogen site location: mixed

H atoms treated by a mixture of independent
and constrained refinement
 $w = 1/[\sigma^2(F_o^2) + (0.020P)^2 + 0.3611P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.48 \text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.44 \text{ e \AA}^{-3}$
Special details

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. Reflections were merged by *SHELXL* according to the crystal class for the calculation of statistics and refinement.

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	0.75812 (2)	0.69658 (2)	0.16087 (2)	0.01737 (4)	
Br2	-0.01570 (2)	0.16791 (2)	0.12324 (2)	0.02003 (5)	
O1	1.18243 (15)	0.98258 (14)	0.33190 (13)	0.0179 (2)	
O2	0.62059 (15)	0.76777 (13)	0.41035 (13)	0.0172 (2)	
H2	0.547 (5)	0.698 (4)	0.323 (3)	0.026*	0.52 (3)
O3	-0.34653 (16)	0.22841 (14)	-0.08661 (15)	0.0227 (2)	
O4	0.33774 (15)	0.50586 (13)	0.19208 (13)	0.0175 (2)	

H4	0.430 (5)	0.603 (3)	0.231 (5)	0.026*	0.48 (3)
O5	0.82422 (16)	0.26379 (15)	0.42261 (14)	0.0219 (2)	
N1	0.48734 (19)	0.29444 (17)	0.32852 (17)	0.0183 (2)	
C1	1.0922 (2)	0.98623 (17)	0.40454 (17)	0.0131 (2)	
C2	0.8905 (2)	0.86761 (17)	0.35306 (17)	0.0138 (2)	
C3	0.7975 (2)	0.87207 (17)	0.44087 (17)	0.0133 (2)	
C4	-0.1856 (2)	0.34979 (18)	-0.04558 (17)	0.0156 (3)	
C5	-0.0041 (2)	0.35843 (17)	0.05629 (17)	0.0149 (3)	
C6	0.1731 (2)	0.49685 (18)	0.10257 (17)	0.0154 (3)	
C7	0.6812 (2)	0.42735 (19)	0.45871 (19)	0.0191 (3)	
H7A	0.6639	0.4839	0.5393	0.023*	
H7B	0.7495	0.5128	0.4135	0.023*	
C8	0.8006 (2)	0.3479 (2)	0.53709 (19)	0.0204 (3)	
H8A	0.9300	0.4358	0.6228	0.025*	
H8B	0.7349	0.2667	0.5868	0.025*	
C9	0.6387 (2)	0.1307 (2)	0.3035 (2)	0.0227 (3)	
H9A	0.5748	0.0515	0.3556	0.027*	
H9B	0.6567	0.0683	0.2278	0.027*	
C10	0.5097 (2)	0.1977 (2)	0.21235 (19)	0.0212 (3)	
H10A	0.5687	0.2716	0.1545	0.025*	
H10B	0.3806	0.1034	0.1327	0.025*	
H1A	0.425 (3)	0.342 (3)	0.278 (3)	0.024 (5)*	
H1B	0.422 (3)	0.224 (3)	0.365 (3)	0.030 (6)*	

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Br1	0.01574 (7)	0.01685 (7)	0.01495 (7)	0.00556 (6)	0.00658 (5)	0.00055 (5)
Br2	0.01706 (8)	0.01455 (7)	0.02405 (8)	0.00653 (6)	0.00668 (6)	0.00865 (6)
O1	0.0161 (5)	0.0199 (5)	0.0195 (5)	0.0081 (4)	0.0110 (4)	0.0050 (4)
O2	0.0108 (5)	0.0164 (5)	0.0184 (5)	0.0030 (4)	0.0064 (4)	0.0015 (4)
O3	0.0136 (5)	0.0158 (5)	0.0308 (6)	0.0039 (4)	0.0068 (5)	0.0081 (4)
O4	0.0128 (5)	0.0138 (5)	0.0195 (5)	0.0051 (4)	0.0035 (4)	0.0038 (4)
O5	0.0167 (5)	0.0258 (6)	0.0240 (6)	0.0130 (5)	0.0079 (4)	0.0038 (5)
N1	0.0146 (6)	0.0210 (6)	0.0244 (7)	0.0108 (5)	0.0105 (5)	0.0118 (5)
C1	0.0124 (6)	0.0127 (6)	0.0150 (6)	0.0069 (5)	0.0061 (5)	0.0057 (5)
C2	0.0124 (6)	0.0134 (6)	0.0134 (6)	0.0056 (5)	0.0051 (5)	0.0025 (5)
C3	0.0113 (6)	0.0125 (6)	0.0151 (6)	0.0064 (5)	0.0046 (5)	0.0049 (5)
C4	0.0168 (7)	0.0131 (6)	0.0161 (6)	0.0063 (5)	0.0083 (5)	0.0028 (5)
C5	0.0156 (6)	0.0123 (6)	0.0161 (6)	0.0067 (5)	0.0071 (5)	0.0047 (5)
C6	0.0166 (7)	0.0139 (6)	0.0139 (6)	0.0060 (5)	0.0076 (5)	0.0021 (5)
C7	0.0195 (7)	0.0172 (7)	0.0223 (7)	0.0095 (6)	0.0107 (6)	0.0059 (6)
C8	0.0187 (7)	0.0251 (7)	0.0177 (7)	0.0129 (6)	0.0066 (6)	0.0051 (6)
C9	0.0209 (7)	0.0185 (7)	0.0255 (8)	0.0108 (6)	0.0073 (6)	0.0028 (6)
C10	0.0185 (7)	0.0206 (7)	0.0189 (7)	0.0093 (6)	0.0047 (6)	0.0022 (6)

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

Br1—C2	1.8807 (14)	C1—C3 ⁱ	1.5282 (19)
Br2—C5	1.8775 (14)	C2—C3	1.3684 (19)
O1—C1	1.2298 (17)	C4—C5	1.446 (2)
O2—C3	1.2937 (17)	C4—C6 ⁱⁱ	1.518 (2)
O2—H2	0.82 (3)	C5—C6	1.363 (2)
O3—C4	1.2231 (18)	C7—C8	1.509 (2)
O4—C6	1.2997 (18)	C7—H7A	0.9900
O4—H4	0.82 (3)	C7—H7B	0.9900
O5—C8	1.4201 (19)	C8—H8A	0.9900
O5—C9	1.428 (2)	C8—H8B	0.9900
N1—C7	1.492 (2)	C9—C10	1.510 (2)
N1—C10	1.493 (2)	C9—H9A	0.9900
N1—H1A	0.86 (3)	C9—H9B	0.9900
N1—H1B	0.85 (3)	C10—H10A	0.9900
C1—C2	1.4377 (19)	C10—H10B	0.9900
C3—O2—H2	118 (3)	C5—C6—C4 ⁱⁱ	119.95 (13)
C6—O4—H4	113 (3)	N1—C7—C8	109.23 (12)
C8—O5—C9	109.77 (12)	N1—C7—H7A	109.8
C7—N1—C10	110.93 (12)	C8—C7—H7A	109.8
C7—N1—H1A	108.5 (14)	N1—C7—H7B	109.8
C10—N1—H1A	108.6 (14)	C8—C7—H7B	109.8
C7—N1—H1B	111.9 (16)	H7A—C7—H7B	108.3
C10—N1—H1B	106.6 (15)	O5—C8—C7	110.62 (13)
H1A—N1—H1B	110 (2)	O5—C8—H8A	109.5
O1—C1—C2	124.48 (13)	C7—C8—H8A	109.5
O1—C1—C3 ⁱ	117.39 (12)	O5—C8—H8B	109.5
C2—C1—C3 ⁱ	118.13 (12)	C7—C8—H8B	109.5
C3—C2—C1	122.62 (13)	H8A—C8—H8B	108.1
C3—C2—Br1	120.30 (11)	O5—C9—C10	111.19 (13)
C1—C2—Br1	117.03 (10)	O5—C9—H9A	109.4
O2—C3—C2	127.87 (13)	C10—C9—H9A	109.4
O2—C3—C1 ⁱ	112.96 (12)	O5—C9—H9B	109.4
C2—C3—C1 ⁱ	119.17 (12)	C10—C9—H9B	109.4
O3—C4—C5	124.23 (13)	H9A—C9—H9B	108.0
O3—C4—C6 ⁱⁱ	118.58 (13)	N1—C10—C9	108.74 (13)
C5—C4—C6 ⁱⁱ	117.19 (12)	N1—C10—H10A	109.9
C6—C5—C4	122.86 (13)	C9—C10—H10A	109.9
C6—C5—Br2	119.20 (11)	N1—C10—H10B	109.9
C4—C5—Br2	117.92 (10)	C9—C10—H10B	109.9
O4—C6—C5	123.68 (13)	H10A—C10—H10B	108.3
O4—C6—C4 ⁱⁱ	116.36 (12)	 	
O1—C1—C2—C3	177.25 (14)	C6 ⁱⁱ —C4—C5—Br2	-178.23 (10)
C3 ⁱ —C1—C2—C3	-3.4 (2)	C4—C5—C6—O4	-178.92 (13)
O1—C1—C2—Br1	-0.21 (19)	Br2—C5—C6—O4	-0.2 (2)

C3 ⁱ —C1—C2—Br1	179.18 (9)	C4—C5—C6—C4 ⁱⁱ	−0.6 (2)
C1—C2—C3—O2	−176.36 (13)	Br2—C5—C6—C4 ⁱⁱ	178.19 (10)
Br1—C2—C3—O2	1.0 (2)	C10—N1—C7—C8	−54.81 (16)
C1—C2—C3—C1 ⁱ	3.4 (2)	C9—O5—C8—C7	−62.38 (16)
Br1—C2—C3—C1 ⁱ	−179.22 (9)	N1—C7—C8—O5	58.53 (17)
O3—C4—C5—C6	−178.65 (15)	C8—O5—C9—C10	62.43 (17)
C6 ⁱⁱ —C4—C5—C6	0.5 (2)	C7—N1—C10—C9	54.27 (17)
O3—C4—C5—Br2	2.6 (2)	O5—C9—C10—N1	−57.89 (18)

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

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

D—H···A	D—H	H···A	D···A	D—H···A
N1—H1A···O4	0.86 (3)	2.04 (3)	2.888 (2)	166 (2)
N1—H1B···O1 ⁱⁱⁱ	0.85 (3)	2.17 (3)	2.938 (2)	151 (2)
N1—H1B···O2 ^{iv}	0.85 (3)	2.29 (3)	2.959 (2)	136 (2)
O2—H2···O4	0.82 (3)	1.77 (3)	2.5174 (16)	153 (4)
O2—H2···O3 ⁱⁱ	0.82 (3)	2.58 (3)	3.0628 (17)	120 (3)
O4—H4···O2	0.82 (3)	1.79 (4)	2.5174 (16)	147 (4)
C7—H7A···O4 ^{iv}	0.99	2.54	3.394 (2)	145
C10—H10B···Br2 ^v	0.99	2.90	3.8905 (17)	175

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

(3)

Crystal data

$\text{C}_4\text{H}_{10}\text{NO}^+\cdot\text{C}_6\text{HBr}_2\text{O}_4^-$	$Z = 2$
$M_r = 385.01$	$F(000) = 376.00$
Triclinic, $P\bar{1}$	$D_x = 2.136 \text{ Mg m}^{-3}$
Hall symbol: -P 1	Mo $K\alpha$ radiation, $\lambda = 0.71075 \text{ \AA}$
$a = 8.62824 (17) \text{ \AA}$	Cell parameters from 15838 reflections
$b = 9.23087 (18) \text{ \AA}$	$\theta = 3.0\text{--}30.1^\circ$
$c = 9.46007 (19) \text{ \AA}$	$\mu = 6.80 \text{ mm}^{-1}$
$\alpha = 93.5321 (7)^\circ$	$T = 180 \text{ K}$
$\beta = 112.9738 (7)^\circ$	Block, brown
$\gamma = 115.9508 (7)^\circ$	$0.40 \times 0.34 \times 0.18 \text{ mm}$
$V = 598.67 (3) \text{ \AA}^3$	

Data collection

Rigaku R-AXIS RAPIDII	3487 independent reflections
diffractometer	3188 reflections with $I > 2\sigma(I)$
Detector resolution: 10.000 pixels mm^{-1}	$R_{\text{int}} = 0.026$
ω scans	$\theta_{\text{max}} = 30.0^\circ$
Absorption correction: numerical	$h = -12 \rightarrow 12$
(NUMABS; Higashi, 1999)	$k = -12 \rightarrow 12$
$T_{\text{min}} = 0.098, T_{\text{max}} = 0.294$	$l = -13 \rightarrow 13$
18199 measured reflections	

Refinement

Refinement on F^2	$R[F^2 > 2\sigma(F^2)] = 0.019$
Least-squares matrix: full	$wR(F^2) = 0.048$

$S = 1.09$
 3487 reflections
 178 parameters
 2 restraints
 Hydrogen site location: mixed

H atoms treated by a mixture of independent
 and constrained refinement
 $w = 1/[\sigma^2(F_o^2) + (0.0216P)^2 + 0.3489P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.001$
 $\Delta\rho_{\text{max}} = 0.59 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.45 \text{ e } \text{\AA}^{-3}$

Special details

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. Reflections were merged by *SHELXL* according to the crystal class for the calculation of statistics and refinement.

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	0.75874 (2)	0.69709 (2)	0.16160 (2)	0.02126 (5)	
Br2	-0.01629 (2)	0.16834 (2)	0.12298 (2)	0.02440 (5)	
O1	1.18225 (16)	0.98340 (14)	0.33207 (13)	0.0212 (2)	
O2	0.62107 (15)	0.76749 (14)	0.41043 (13)	0.0205 (2)	
H2	0.550 (6)	0.702 (5)	0.321 (3)	0.031*	0.50 (3)
O3	-0.34638 (16)	0.22923 (14)	-0.08655 (16)	0.0275 (2)	
O4	0.33726 (16)	0.50565 (14)	0.19202 (13)	0.0209 (2)	
H4	0.426 (5)	0.604 (3)	0.231 (5)	0.031*	0.50 (3)
O5	0.82325 (17)	0.26349 (16)	0.42251 (15)	0.0266 (2)	
N1	0.48719 (19)	0.29464 (18)	0.32897 (17)	0.0216 (3)	
C1	1.0921 (2)	0.98669 (17)	0.40464 (17)	0.0155 (2)	
C2	0.8904 (2)	0.86771 (17)	0.35338 (17)	0.0158 (2)	
C3	0.7977 (2)	0.87190 (17)	0.44099 (17)	0.0157 (2)	
C4	-0.1853 (2)	0.35010 (18)	-0.04556 (18)	0.0185 (3)	
C5	-0.0045 (2)	0.35849 (17)	0.05607 (17)	0.0175 (3)	
C6	0.1728 (2)	0.49667 (18)	0.10245 (17)	0.0180 (3)	
C7	0.6809 (2)	0.42714 (19)	0.4585 (2)	0.0227 (3)	
H7A	0.6639	0.4840	0.5389	0.027*	
H7B	0.7492	0.5122	0.4133	0.027*	
C8	0.7997 (2)	0.3474 (2)	0.53683 (19)	0.0250 (3)	
H8A	0.9291	0.4350	0.6226	0.030*	
H8B	0.7337	0.2662	0.5861	0.030*	
C9	0.6381 (2)	0.1310 (2)	0.3036 (2)	0.0272 (3)	
H9A	0.5742	0.0517	0.3552	0.033*	
H9B	0.6562	0.0690	0.2279	0.033*	
C10	0.5094 (2)	0.1980 (2)	0.21303 (19)	0.0252 (3)	
H10A	0.5683	0.2719	0.1554	0.030*	
H10B	0.3803	0.1040	0.1335	0.030*	
H1A	0.424 (3)	0.344 (3)	0.277 (2)	0.025 (5)*	
H1B	0.422 (3)	0.223 (3)	0.367 (3)	0.026 (5)*	

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Br1	0.01905 (8)	0.02063 (8)	0.01805 (7)	0.00654 (6)	0.00783 (6)	0.00043 (5)
Br2	0.02061 (8)	0.01771 (7)	0.02936 (9)	0.00800 (6)	0.00791 (6)	0.01079 (6)
O1	0.0192 (5)	0.0234 (5)	0.0227 (5)	0.0093 (4)	0.0131 (4)	0.0057 (4)
O2	0.0128 (5)	0.0201 (5)	0.0215 (5)	0.0035 (4)	0.0076 (4)	0.0019 (4)
O3	0.0161 (5)	0.0194 (5)	0.0374 (7)	0.0049 (4)	0.0080 (5)	0.0101 (5)
O4	0.0151 (5)	0.0166 (5)	0.0230 (5)	0.0061 (4)	0.0039 (4)	0.0046 (4)
O5	0.0192 (5)	0.0314 (6)	0.0295 (6)	0.0151 (5)	0.0094 (5)	0.0045 (5)
N1	0.0171 (6)	0.0245 (6)	0.0290 (7)	0.0129 (5)	0.0123 (5)	0.0141 (6)
C1	0.0148 (6)	0.0147 (6)	0.0180 (6)	0.0076 (5)	0.0079 (5)	0.0069 (5)
C2	0.0141 (6)	0.0144 (6)	0.0159 (6)	0.0057 (5)	0.0062 (5)	0.0027 (5)
C3	0.0133 (6)	0.0155 (6)	0.0180 (6)	0.0076 (5)	0.0064 (5)	0.0061 (5)
C4	0.0181 (7)	0.0155 (6)	0.0197 (6)	0.0065 (5)	0.0092 (5)	0.0034 (5)
C5	0.0175 (6)	0.0141 (6)	0.0192 (6)	0.0073 (5)	0.0078 (5)	0.0053 (5)
C6	0.0188 (7)	0.0164 (6)	0.0166 (6)	0.0071 (5)	0.0086 (5)	0.0030 (5)
C7	0.0231 (7)	0.0208 (7)	0.0259 (7)	0.0111 (6)	0.0129 (6)	0.0066 (6)
C8	0.0219 (7)	0.0308 (8)	0.0218 (7)	0.0150 (7)	0.0080 (6)	0.0061 (6)
C9	0.0250 (8)	0.0217 (7)	0.0317 (8)	0.0129 (7)	0.0095 (7)	0.0036 (6)
C10	0.0218 (7)	0.0243 (7)	0.0220 (7)	0.0108 (6)	0.0051 (6)	0.0025 (6)

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

Br1—C2	1.8803 (14)	C1—C3 ⁱ	1.5283 (19)
Br2—C5	1.8769 (14)	C2—C3	1.367 (2)
O1—C1	1.2299 (17)	C4—C5	1.442 (2)
O2—C3	1.2931 (17)	C4—C6 ⁱⁱ	1.519 (2)
O2—H2	0.82 (3)	C5—C6	1.363 (2)
O3—C4	1.2229 (18)	C7—C8	1.509 (2)
O4—C6	1.2997 (18)	C7—H7A	0.9900
O4—H4	0.82 (3)	C7—H7B	0.9900
O5—C8	1.420 (2)	C8—H8A	0.9900
O5—C9	1.425 (2)	C8—H8B	0.9900
N1—C7	1.490 (2)	C9—C10	1.508 (2)
N1—C10	1.493 (2)	C9—H9A	0.9900
N1—H1A	0.89 (3)	C9—H9B	0.9900
N1—H1B	0.86 (3)	C10—H10A	0.9900
C1—C2	1.4394 (19)	C10—H10B	0.9900
C3—O2—H2	117 (3)	C5—C6—C4 ⁱⁱ	119.94 (13)
C6—O4—H4	111 (3)	N1—C7—C8	109.14 (13)
C8—O5—C9	109.83 (12)	N1—C7—H7A	109.9
C7—N1—C10	110.85 (12)	C8—C7—H7A	109.9
C7—N1—H1A	108.4 (14)	N1—C7—H7B	109.9
C10—N1—H1A	108.2 (14)	C8—C7—H7B	109.9
C7—N1—H1B	111.2 (14)	H7A—C7—H7B	108.3
C10—N1—H1B	106.4 (14)	O5—C8—C7	110.51 (13)

H1A—N1—H1B	112 (2)	O5—C8—H8A	109.5
O1—C1—C2	124.51 (13)	C7—C8—H8A	109.5
O1—C1—C3 ⁱ	117.41 (12)	O5—C8—H8B	109.5
C2—C1—C3 ⁱ	118.08 (12)	C7—C8—H8B	109.5
C3—C2—C1	122.63 (13)	H8A—C8—H8B	108.1
C3—C2—Br1	120.43 (11)	O5—C9—C10	111.18 (13)
C1—C2—Br1	116.90 (10)	O5—C9—H9A	109.4
O2—C3—C2	127.81 (13)	C10—C9—H9A	109.4
O2—C3—C1 ⁱ	112.98 (12)	O5—C9—H9B	109.4
C2—C3—C1 ⁱ	119.21 (12)	C10—C9—H9B	109.4
O3—C4—C5	124.27 (14)	H9A—C9—H9B	108.0
O3—C4—C6 ⁱⁱ	118.48 (14)	N1—C10—C9	108.84 (13)
C5—C4—C6 ⁱⁱ	117.24 (12)	N1—C10—H10A	109.9
C6—C5—C4	122.82 (13)	C9—C10—H10A	109.9
C6—C5—Br2	119.20 (11)	N1—C10—H10B	109.9
C4—C5—Br2	117.98 (10)	C9—C10—H10B	109.9
O4—C6—C5	123.69 (14)	H10A—C10—H10B	108.3
O4—C6—C4 ⁱⁱ	116.35 (13)		
O1—C1—C2—C3	177.30 (14)	C6 ⁱⁱ —C4—C5—Br2	-178.27 (10)
C3 ⁱ —C1—C2—C3	-3.3 (2)	C4—C5—C6—O4	-178.95 (14)
O1—C1—C2—Br1	-0.20 (19)	Br2—C5—C6—O4	-0.2 (2)
C3 ⁱ —C1—C2—Br1	179.24 (9)	C4—C5—C6—C4 ⁱⁱ	-0.5 (2)
C1—C2—C3—O2	-176.48 (14)	Br2—C5—C6—C4 ⁱⁱ	178.25 (10)
Br1—C2—C3—O2	0.9 (2)	C10—N1—C7—C8	-55.03 (17)
C1—C2—C3—C1 ⁱ	3.3 (2)	C9—O5—C8—C7	-62.46 (17)
Br1—C2—C3—C1 ⁱ	-179.29 (9)	N1—C7—C8—O5	58.77 (17)
O3—C4—C5—C6	-178.51 (15)	C8—O5—C9—C10	62.31 (18)
C6 ⁱⁱ —C4—C5—C6	0.5 (2)	C7—N1—C10—C9	54.35 (17)
O3—C4—C5—Br2	2.7 (2)	O5—C9—C10—N1	-57.76 (19)

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

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

$D\text{—H}\cdots A$	$D\text{—H}$	$H\cdots A$	$D\cdots A$	$D\text{—H}\cdots A$
N1—H1A…O4	0.89 (3)	2.02 (3)	2.890 (2)	167 (2)
N1—H1B…O1 ⁱⁱⁱ	0.86 (3)	2.16 (3)	2.938 (2)	150 (2)
N1—H1B…O2 ^{iv}	0.86 (3)	2.28 (3)	2.964 (2)	136 (2)
O2—H2…O4	0.82 (3)	1.79 (4)	2.5224 (16)	148 (5)
O2—H2…O3 ⁱⁱ	0.82 (3)	2.55 (4)	3.0678 (18)	122 (4)
O4—H4…O2	0.82 (3)	1.80 (4)	2.5224 (16)	147 (4)
C7—H7A…O4 ^{iv}	0.99	2.55	3.402 (2)	145
C10—H10B…Br2 ^v	0.99	2.91	3.8946 (17)	174

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