

## N-(4-Bromophenyl)propionamide

Chen-Hu Guo, Ming-Lin Guo,\* Hou-Ying Zhang and Ya-Nan Ye

School of Materials and Chemical Engineering, and Key Laboratory of Hollow Fiber Membrane Materials and Membrane Processes, Tianjin Polytechnic University, Tianjin 300160, People's Republic of China

Correspondence e-mail: guomlin@yahoo.com

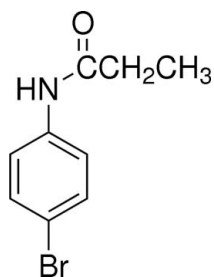
Received 17 August 2007; accepted 19 August 2007

Key indicators: single-crystal X-ray study;  $T = 294$  K; mean  $\sigma(\text{C}-\text{C}) = 0.004$  Å;  $R$  factor = 0.036;  $wR$  factor = 0.093; data-to-parameter ratio = 14.4.

In the title compound,  $\text{C}_9\text{H}_{10}\text{BrNO}$ , molecules are interconnected by a framework of intermolecular  $\text{N}-\text{H}\cdots\text{O}$  hydrogen bonds. Infinite one-dimensional hydrogen-bonded chains extend along the  $a$  direction. These chains are aggregated into layers through weak inter-chain  $\text{C}-\text{H}\cdots\text{Br}$  contacts along  $c$ .

### Related literature

For structural studies on other amides, see: Knopfel *et al.* (2005); Guo (2004); Parvez *et al.* (2004). For bond-length data, see: Allen *et al.* (1987).



### Experimental

#### Crystal data

$\text{C}_9\text{H}_{10}\text{BrNO}$   
 $M_r = 228.09$

Orthorhombic,  $Pbca$   
 $a = 9.3879$  (19) Å

$b = 9.2952$  (19) Å  
 $c = 20.721$  (4) Å  
 $V = 1808.2$  (6) Å<sup>3</sup>  
 $Z = 8$

Mo  $K\alpha$  radiation  
 $\mu = 4.50$  mm<sup>-1</sup>  
 $T = 294$  (2) K  
 $0.08 \times 0.08 \times 0.04$  mm

#### Data collection

Rigaku Saturn diffractometer  
Absorption correction: multi-scan  
(Jacobson, 1998)  
 $T_{\min} = 0.712$ ,  $T_{\max} = 0.835$

10043 measured reflections  
1587 independent reflections  
1375 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.064$

#### Refinement

$R[F^2 > 2\sigma(F^2)] = 0.036$   
 $wR(F^2) = 0.093$   
 $S = 1.10$   
1587 reflections

110 parameters  
H-atom parameters constrained  
 $\Delta\rho_{\max} = 0.59$  e Å<sup>-3</sup>  
 $\Delta\rho_{\min} = -0.76$  e Å<sup>-3</sup>

**Table 1**

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{C9}-\text{H9A}\cdots\text{Br1}^{\text{i}}$	0.96	2.99	3.555 (3)	119
$\text{C2}-\text{H2}\cdots\text{O1}$	0.93	2.32	2.880 (4)	118
$\text{N1}-\text{H1A}\cdots\text{O1}^{\text{ii}}$	0.85	2.08	2.886 (3)	160

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

Data collection: *CrystalClear* (Rigaku/MS, 2005); cell refinement: *CrystalClear*; data reduction: *CrystalClear*; program(s) used to solve structure: *SHELXTL* (Bruker, 2001); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

We thank Tianjin Polytechnic University for financial support.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: SJ2337).

### 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.
- Bruker (2001). *SHELXTL*. Version 6.12. Bruker AXS Inc., Madison, Wisconsin, USA.
- Guo, M.-L. (2004). *Acta Cryst. E60*, o736–o737.
- Jacobson, R. (1998). Private communication to the Rigaku Corporation, Tokyo, Japan.
- Knopfel, T. F., Zarotti, P., Ichikawa, T. & Carreira, E. M. (2005). *J. Am. Chem. Soc.* **127**, 9682–9683.
- Parvez, M., Shahid, K., Shahzadi, S. & Ali, S. (2004). *Acta Cryst. E60*, o2079–o2081.
- Rigaku/MS (2005). *CrystalClear*. Version 1.3.6. Rigaku/MS, The Woodlands, Texas, USA.

**supplementary materials**

*Acta Cryst.* (2007). E63, o3884 [ doi:10.1107/S1600536807040950 ]

## *N*-(4-Bromophenyl)propionamide

C.-H. Guo, M.-L. Guo, H.-Y. Zhang and Y.-N. Ye

### Comment

The amide moiety is an important constituent of many biologically significant compounds. Structural studies of amides are therefore of interest (Knopfel *et al.*, 2005; Guo, 2004; Parvez *et al.*, 2004). As part of a study of possible drugs, the crystal structure of the title compound (I) is reported herein, Fig. 1. Bond distances and angles are normal, within experimental error (Allen *et al.*, 1987).

In the structure of (I), the molecular skeleton is essentially planar with atoms O1, C7, C8 and C9 lying 0.6630, 0.2724, 0.0990 and 0.1530 Å, respectively, out of the N1/C1—C6/Br1 mean plane. Intermolecular N1—H1A $\cdots$ O1<sup>i</sup> hydrogen bonds (see Table 1 for symmetry code) link the molecules into chains (Fig. 2). Further, short C—H $\cdots$ Br contacts (distance = 3.555 (3) Å) stabilize the packing in the structure.

### Experimental

The title compound was prepared by slowly adding concentrated sulfuric acid (2.0 ml) dropwise to a mixture of *N*-phenylpropionamide (3.7 g, 0.025 mol), sodium bromide (2.6 g, 0.025 mol), 30% hydrogen peroxide (5.0 ml, 0.05 mol) and ethanol (20 ml). The mixture was stirred in air at room temperature for 30 min. The resulting product was separated by filtration (yield 5.3 g, 93%). Single crystals were obtained from a solution of 0.2 g of the product in 25 ml of distilled water by slow concentration over a period of 1 week at room temperature.

### Refinement

All H-atoms were positioned geometrically and refined using a riding model with  $d(\text{C—H}) = 0.93$  Å,  $U_{\text{iso}} = 1.2U_{\text{eq}}(\text{C})$  for aromatic, 0.97 Å,  $U_{\text{iso}} = 1.2U_{\text{eq}}(\text{C})$  for CH<sub>2</sub>, 0.96 Å,  $U_{\text{iso}} = 1.5U_{\text{eq}}(\text{C})$  for CH<sub>3</sub> atoms and 0.86 Å,  $U_{\text{iso}} = 1.2U_{\text{eq}}(\text{N})$  for the NH atom.

### Figures

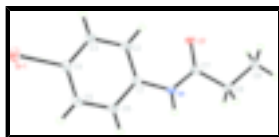


Fig. 1. A view of the structure of (I), showing the atom-numbering Scheme; displacement ellipsoids were drawn at the 30% probability level.

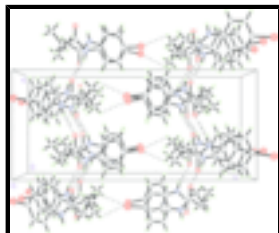


Fig. 2. The crystal packing of (I) viewed down the *b* axis with hydrogen bonds drawn as dashed lines.

## *N*-(4-Bromophenyl)propionamide

### *Crystal data*

$C_9H_{10}BrNO$	$F_{000} = 912$
$M_r = 228.09$	$D_x = 1.676 \text{ Mg m}^{-3}$
Orthorhombic, <i>Pbca</i>	Mo $K\alpha$ radiation
Hall symbol: -P 2ac 2ab	$\lambda = 0.71073 \text{ \AA}$
$a = 9.3879 (19) \text{ \AA}$	Cell parameters from 3850 reflections
$b = 9.2952 (19) \text{ \AA}$	$\theta = 2.0\text{--}27.9^\circ$
$c = 20.721 (4) \text{ \AA}$	$\mu = 4.50 \text{ mm}^{-1}$
$V = 1808.2 (6) \text{ \AA}^3$	$T = 294 (2) \text{ K}$
$Z = 8$	Plate, colorless
	$0.08 \times 0.08 \times 0.04 \text{ mm}$

### *Data collection*

Rigaku Saturn diffractometer	1587 independent reflections
Radiation source: rotating anode	1375 reflections with $I > 2\sigma(I)$
Monochromator: confocal	$R_{\text{int}} = 0.064$
$T = 294(2) \text{ K}$	$\theta_{\text{max}} = 25.0^\circ$
$\omega$ scans	$\theta_{\text{min}} = 2.0^\circ$
Absorption correction: multi-scan (Jacobson, 1998)	$h = -11 \rightarrow 7$
$T_{\text{min}} = 0.712$ , $T_{\text{max}} = 0.835$	$k = -10 \rightarrow 11$
10043 measured reflections	$l = -24 \rightarrow 24$

### *Refinement*

Refinement on $F^2$	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.036$	H-atom parameters constrained
$wR(F^2) = 0.093$	$w = 1/[\sigma^2(F_o^2) + (0.0501P)^2]$
$S = 1.10$	where $P = (F_o^2 + 2F_c^2)/3$
1587 reflections	$(\Delta/\sigma)_{\text{max}} = 0.001$
110 parameters	$\Delta\rho_{\text{max}} = 0.59 \text{ e \AA}^{-3}$
Primary atom site location: structure-invariant direct methods	$\Delta\rho_{\text{min}} = -0.75 \text{ e \AA}^{-3}$
	Extinction correction: none

### *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.** 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 ( $\text{\AA}^2$ )

	$x$	$y$	$z$	$U_{\text{iso}}^*/U_{\text{eq}}$
Br1	0.23826 (4)	0.03223 (3)	0.528300 (15)	0.03307 (18)
N1	0.3167 (3)	0.4370 (2)	0.30106 (11)	0.0211 (5)
H1A	0.4042	0.4501	0.2932	0.025*
O1	0.0960 (2)	0.45235 (19)	0.25674 (9)	0.0275 (5)
C1	0.2921 (3)	0.3403 (3)	0.35205 (13)	0.0195 (6)
C2	0.1577 (3)	0.3134 (3)	0.37778 (14)	0.0245 (7)
H2	0.0780	0.3580	0.3602	0.029*
C3	0.1428 (3)	0.2203 (3)	0.42943 (14)	0.0248 (7)
H3	0.0530	0.2016	0.4464	0.030*
C4	0.2612 (3)	0.1554 (3)	0.45556 (15)	0.0238 (7)
C5	0.3955 (3)	0.1803 (3)	0.43096 (14)	0.0280 (7)
H5	0.4746	0.1353	0.4489	0.034*
C6	0.4107 (3)	0.2730 (3)	0.37931 (14)	0.0251 (7)
H6	0.5009	0.2908	0.3625	0.030*
C7	0.2219 (3)	0.4876 (3)	0.25727 (16)	0.0209 (7)
C8	0.2844 (3)	0.5911 (3)	0.20894 (15)	0.0238 (7)
H8A	0.3453	0.5384	0.1794	0.029*
H8B	0.3432	0.6607	0.2315	0.029*
C9	0.1713 (3)	0.6702 (3)	0.17065 (15)	0.0294 (7)
H9A	0.1148	0.6021	0.1471	0.044*
H9B	0.2162	0.7353	0.1410	0.044*
H9C	0.1113	0.7234	0.1996	0.044*

Atomic displacement parameters ( $\text{\AA}^2$ )

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
Br1	0.0451 (3)	0.0296 (3)	0.0246 (3)	0.00330 (13)	0.00111 (13)	0.00321 (12)
N1	0.0113 (12)	0.0276 (12)	0.0244 (13)	-0.0007 (10)	0.0004 (11)	0.0004 (10)
O1	0.0147 (11)	0.0365 (11)	0.0314 (12)	-0.0016 (9)	-0.0024 (9)	0.0040 (9)
C1	0.0200 (15)	0.0187 (14)	0.0200 (15)	0.0013 (12)	-0.0006 (12)	-0.0058 (12)
C2	0.0153 (15)	0.0315 (15)	0.0267 (17)	-0.0013 (12)	-0.0021 (13)	-0.0026 (13)
C3	0.0210 (16)	0.0309 (15)	0.0224 (17)	-0.0044 (13)	-0.0007 (13)	-0.0025 (12)
C4	0.0333 (18)	0.0202 (15)	0.0177 (15)	0.0001 (12)	-0.0009 (13)	-0.0006 (14)
C5	0.0245 (17)	0.0319 (15)	0.0276 (17)	0.0077 (14)	-0.0042 (14)	-0.0024 (14)
C6	0.0180 (15)	0.0304 (15)	0.0269 (17)	0.0034 (12)	0.0007 (13)	-0.0014 (13)
C7	0.0168 (16)	0.0246 (14)	0.0212 (17)	0.0019 (12)	0.0012 (12)	-0.0060 (12)
C8	0.0202 (16)	0.0249 (15)	0.0263 (17)	-0.0023 (12)	0.0014 (12)	0.0032 (14)

# supplementary materials

C9                    0.0282 (17)            0.0292 (15)            0.0308 (18)            0.0024 (14)            0.0007 (15)            0.0041 (14)

## Geometric parameters (Å, °)

Br1—C4	1.905 (3)	C4—C5	1.380 (4)
N1—C7	1.355 (4)	C5—C6	1.382 (4)
N1—C1	1.406 (4)	C5—H5	0.9300
N1—H1A	0.8457	C6—H6	0.9300
O1—C7	1.226 (3)	C7—C8	1.508 (4)
C1—C2	1.392 (4)	C8—C9	1.516 (4)
C1—C6	1.397 (4)	C8—H8A	0.9700
C2—C3	1.383 (4)	C8—H8B	0.9700
C2—H2	0.9300	C9—H9A	0.9600
C3—C4	1.376 (4)	C9—H9B	0.9600
C3—H3	0.9300	C9—H9C	0.9600
C7—N1—C1	128.1 (3)	C5—C6—C1	120.7 (3)
C7—N1—H1A	117.3	C5—C6—H6	119.7
C1—N1—H1A	113.3	C1—C6—H6	119.7
C2—C1—C6	119.1 (3)	O1—C7—N1	123.1 (3)
C2—C1—N1	123.5 (3)	O1—C7—C8	122.6 (3)
C6—C1—N1	117.3 (3)	N1—C7—C8	114.2 (3)
C3—C2—C1	120.1 (3)	C7—C8—C9	112.6 (3)
C3—C2—H2	120.0	C7—C8—H8A	109.1
C1—C2—H2	120.0	C9—C8—H8A	109.1
C4—C3—C2	119.8 (3)	C7—C8—H8B	109.1
C4—C3—H3	120.1	C9—C8—H8B	109.1
C2—C3—H3	120.1	H8A—C8—H8B	107.8
C3—C4—C5	121.3 (3)	C8—C9—H9A	109.5
C3—C4—Br1	118.9 (2)	C8—C9—H9B	109.5
C5—C4—Br1	119.8 (2)	H9A—C9—H9B	109.5
C4—C5—C6	119.0 (3)	C8—C9—H9C	109.5
C4—C5—H5	120.5	H9A—C9—H9C	109.5
C6—C5—H5	120.5	H9B—C9—H9C	109.5

## Hydrogen-bond geometry (Å, °)

<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>
C9—H9A $\cdots$ Br1 <sup>i</sup>	0.96	2.99	3.555 (3)	119
C2—H2 $\cdots$ O1	0.93	2.32	2.880 (4)	118
N1—H1A $\cdots$ O1 <sup>ii</sup>	0.85	2.08	2.886 (3)	160

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

Fig. 1

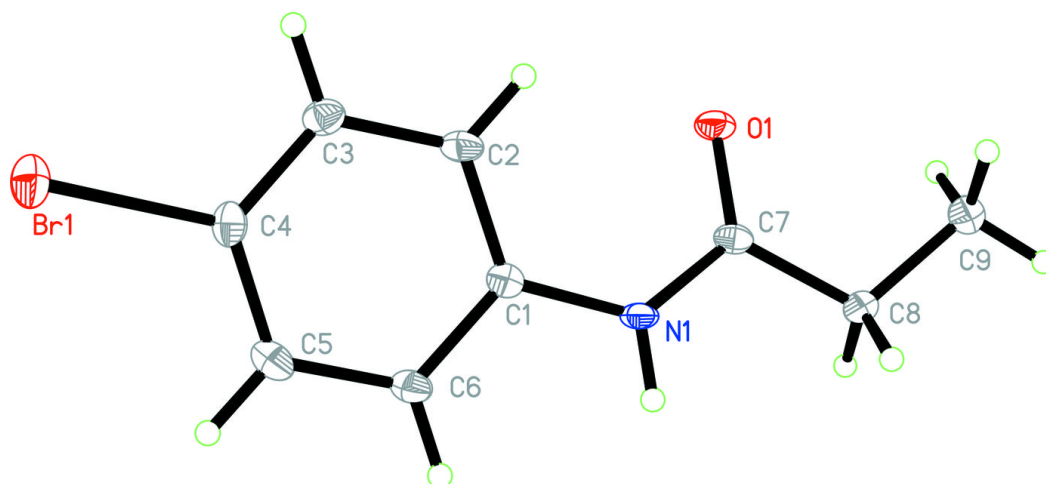


Fig. 2

