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## Structure Reports

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## 2-(4-Bromophenyl)-2-methylpropanamide

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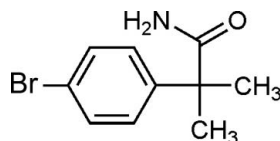
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 Key indicators: single-crystal X-ray study;  $T = 113$  K; mean  $\sigma(\text{C}-\text{C}) = 0.004$  Å;  $R$  factor = 0.035;  $wR$  factor = 0.074; data-to-parameter ratio = 14.0.

In the crystal of the title compound,  $\text{C}_{10}\text{H}_{12}\text{BrNO}$ , inversion dimers linked by pairs of  $\text{N}-\text{H}\cdots\text{O}$  hydrogen bonds generate  $R_2^2(8)$  loops. Further  $\text{N}-\text{H}\cdots\text{O}$  hydrogen bonds link the dimers into sheets propagating in (100).

### Related literature

 For the synthesis, see: Koltunov *et al.* (2004).


### Experimental

#### Crystal data

$\text{C}_{10}\text{H}_{12}\text{BrNO}$   
 $M_r = 242.12$   
 Monoclinic,  $P2_1/c$   
 $a = 16.425$  (8) Å  
 $b = 6.135$  (3) Å  
 $c = 10.152$  (5) Å

$\beta = 97.613$  (7)°  
 $V = 1013.9$  (8) Å<sup>3</sup>  
 $Z = 4$   
 Mo  $K\alpha$  radiation  
 $\mu = 4.01$  mm<sup>-1</sup>  
 $T = 113$  K

 $0.20 \times 0.18 \times 0.12$  mm

#### Data collection

Rigaku Saturn CCD diffractometer  
 Absorption correction: multi-scan  
 (*CrystalClear*; Rigaku/MS, 2005)  
 $T_{\min} = 0.501$ ,  $T_{\max} = 0.644$

9829 measured reflections  
 1787 independent reflections  
 1333 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.090$

#### Refinement

$R[F^2 > 2\sigma(F^2)] = 0.035$   
 $wR(F^2) = 0.074$   
 $S = 0.99$   
 1787 reflections  
 128 parameters  
 3 restraints

H atoms treated by a mixture of independent and constrained refinement  
 $\Delta\rho_{\max} = 0.55$  e Å<sup>-3</sup>  
 $\Delta\rho_{\min} = -0.53$  e Å<sup>-3</sup>

**Table 1**

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N1}-\text{H1A}\cdots\text{O1}^{\text{i}}$	0.89 (1)	2.12 (1)	2.990 (3)	167 (3)
$\text{N1}-\text{H1B}\cdots\text{O1}^{\text{ii}}$	0.88 (1)	2.12 (1)	3.002 (3)	173 (3)

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

Data collection: *CrystalClear* (Rigaku/MS, 2005); cell refinement: *CrystalClear*; data reduction: *CrystalClear*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *CrystalStructure* (Rigaku/MS, 2005).

We thank the College Research Program of Yuncheng University (2008113) for funding.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB5360).

### References

- Koltunov, K. Y., Walspurger, S. & Sommer, J. (2004). *Eur. J. Org. Chem.* pp. 4039–4047.  
 Rigaku/MS (2005). *CrystalClear* and *CrystalStructure*. Rigaku/MS Inc., The Woodlands, Texas, USA.  
 Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112–122.

## supporting information

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**S1. Comment**

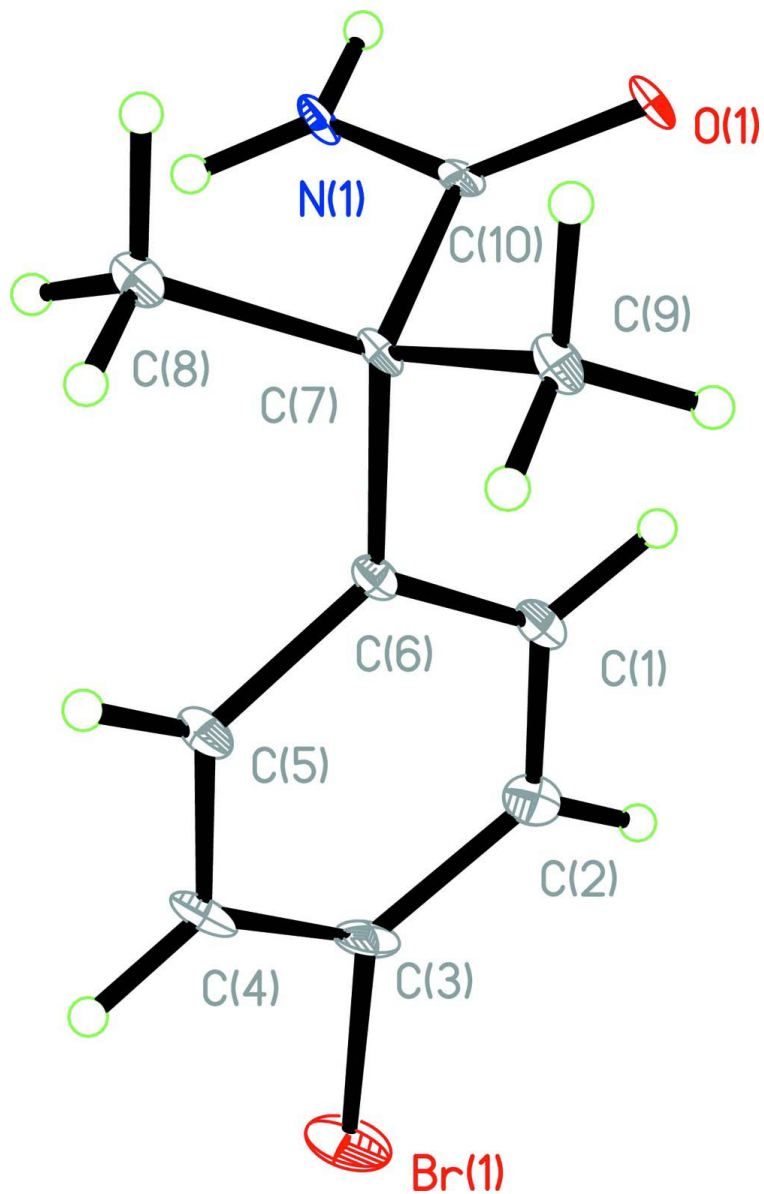
The reaction of amides towards weak nucleophiles such as nonactivated arenes have very broad utility in organic chemistry. However, little work has been done to investigate it. The title compound was synthesized by a facile method through the reaction of methacrylamide and benzene, catalyzed by  $\text{AlCl}_3$ . In the crystal, the molecules are linked by intermolecular N—H $\cdots$ O hydrogen bonding interactions. Single-crystal X-ray diffraction analysis reveals that the title compound crystallizes in the Monoclinic space group  $P 21/c$ .

**S2. Experimental**

A mixture of  $\text{AlCl}_3$  (0.95 g, 7.1 mmol) and methacrylamide (0.2 g, 2.3 mmol) in benzene (3 ml) was stirred at 25 °C for 3 h, and was then poured over several grams of ice and extracted with  $\text{CH}_2\text{Cl}_2$ . The organic phase was separated, dried with anhydrous  $\text{Na}_2\text{SO}_4$  and concentrated in vacuo to give a solid mixture of 2-methyl-3-phenylpropionamide and 2-methyl-2-phenylpropionamide (0.41 g, 97%) in 2:1 ratio. The title compound was separated by flash column chromatography on silica gel. Colourless prisms of (I) were obtained by recrystallisation from ethanol.

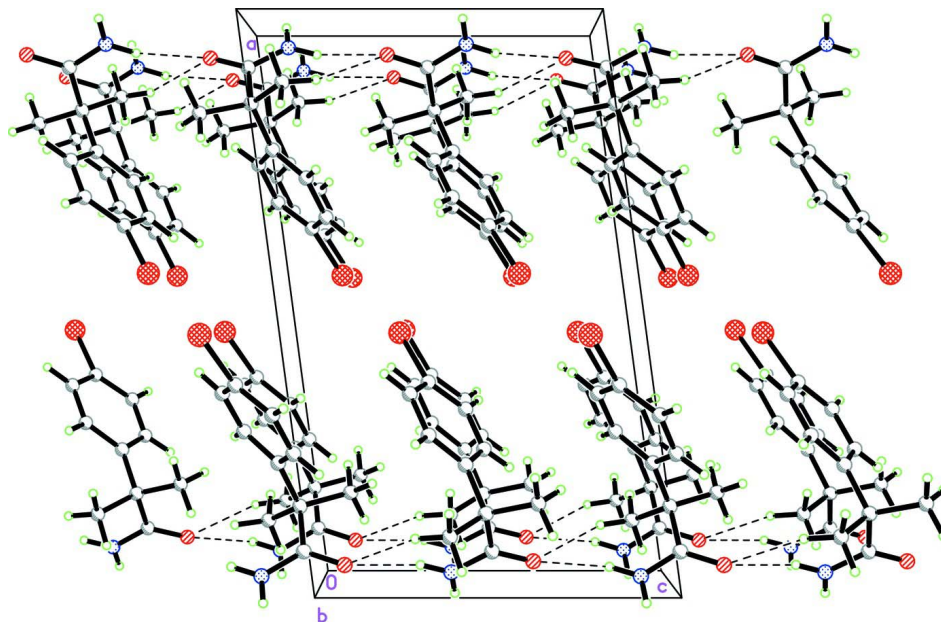
**S3. Refinement**

The H atoms were positioned geometrically (C—H=0.95 Å or 0.98 Å, N—H=0.88 Å) and refined as riding with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C}, \text{N})$  or  $1.5U_{\text{eq}}(\text{methyl C})$ .



**Figure 1**

The molecular structure of (I). Displacement ellipsoids are drawn at the 30% probability level and H atoms are shown as spheres of arbitrary radius.

**Figure 2**

The crystal packing for (I).

### 2-(4-Bromophenyl)-2-methylpropanamide

#### Crystal data

$C_{10}H_{12}BrNO$

$M_r = 242.12$

Monoclinic,  $P2_1/c$

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

$a = 16.425\ (8)\ \text{\AA}$

$b = 6.135\ (3)\ \text{\AA}$

$c = 10.152\ (5)\ \text{\AA}$

$\beta = 97.613\ (7)^\circ$

$V = 1013.9\ (8)\ \text{\AA}^3$

$Z = 4$

$F(000) = 488$

$D_x = 1.586\ \text{Mg m}^{-3}$

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

Cell parameters from 3466 reflections

$\theta = 2.2\text{--}27.9^\circ$

$\mu = 4.01\ \text{mm}^{-1}$

$T = 113\ \text{K}$

Prism, colourless

$0.20 \times 0.18 \times 0.12\ \text{mm}$

#### Data collection

Rigaku Saturn CCD  
diffractometer

Radiation source: rotating anode

Multilayer monochromator

Detector resolution:  $14.63\ \text{pixels mm}^{-1}$

$\omega$  and  $\varphi$  scans

Absorption correction: multi-scan

(*CrystalClear*; Rigaku/MSO, 2005)

$T_{\min} = 0.501$ ,  $T_{\max} = 0.644$

9829 measured reflections

1787 independent reflections

1333 reflections with  $I > 2\sigma(I)$

$R_{\text{int}} = 0.090$

$\theta_{\max} = 25.0^\circ$ ,  $\theta_{\min} = 2.5^\circ$

$h = -19 \rightarrow 19$

$k = -7 \rightarrow 7$

$l = -12 \rightarrow 12$

#### Refinement

Refinement on  $F^2$

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.035$

$wR(F^2) = 0.074$

$S = 0.99$

1787 reflections

128 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 atoms treated by a mixture of independent and constrained refinement

$$w = 1/[\sigma^2(F_o^2) + (0.0242P)^2]$$

where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} = 0.001$   
 $\Delta\rho_{\max} = 0.55 \text{ e } \text{\AA}^{-3}$   
 $\Delta\rho_{\min} = -0.53 \text{ e } \text{\AA}^{-3}$

*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  $F^2 > \sigma(F^2)$  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 ( $\text{\AA}^2$ )*

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
Br1	0.44964 (2)	0.60527 (5)	0.82615 (4)	0.03800 (16)
O1	0.07026 (12)	0.3192 (3)	1.10663 (18)	0.0161 (5)
N1	0.05185 (16)	0.3190 (4)	0.8836 (2)	0.0152 (6)
C1	0.25495 (19)	0.4020 (4)	1.0149 (3)	0.0172 (7)
H1	0.2230	0.4624	1.0773	0.021*
C2	0.3202 (2)	0.5183 (5)	0.9799 (3)	0.0211 (7)
H2	0.3331	0.6576	1.0180	0.025*
C3	0.36720 (19)	0.4317 (5)	0.8887 (3)	0.0207 (7)
C4	0.35071 (19)	0.2238 (5)	0.8383 (3)	0.0241 (8)
H4	0.3844	0.1611	0.7792	0.029*
C5	0.28481 (19)	0.1088 (5)	0.8747 (3)	0.0178 (7)
H5	0.2737	-0.0335	0.8400	0.021*
C6	0.23442 (18)	0.1968 (4)	0.9608 (3)	0.0124 (6)
C7	0.15824 (18)	0.0770 (4)	0.9971 (3)	0.0122 (7)
C8	0.12810 (19)	-0.1034 (4)	0.8969 (3)	0.0170 (7)
H8A	0.1223	-0.0439	0.8066	0.026*
H8B	0.0748	-0.1582	0.9159	0.026*
H8C	0.1680	-0.2230	0.9044	0.026*
C9	0.17838 (19)	-0.0249 (4)	1.1358 (3)	0.0168 (7)
H9A	0.2222	-0.1331	1.1346	0.025*
H9B	0.1292	-0.0960	1.1608	0.025*
H9C	0.1966	0.0893	1.2005	0.025*
C10	0.08953 (17)	0.2487 (4)	1.0001 (3)	0.0117 (6)
H1A	0.0657 (16)	0.283 (4)	0.8050 (15)	0.025 (9)*
H1B	0.0127 (13)	0.417 (4)	0.885 (2)	0.025 (9)*

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

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
Br1	0.0253 (3)	0.0389 (3)	0.0549 (3)	-0.00663 (17)	0.0241 (2)	0.00685 (18)

O1	0.0196 (13)	0.0189 (11)	0.0122 (11)	0.0030 (9)	0.0113 (9)	-0.0008 (9)
N1	0.0169 (15)	0.0182 (14)	0.0121 (14)	0.0065 (12)	0.0079 (12)	-0.0018 (11)
C1	0.0176 (19)	0.0195 (17)	0.0161 (16)	0.0028 (14)	0.0078 (14)	0.0009 (13)
C2	0.0189 (19)	0.0212 (17)	0.0237 (18)	0.0021 (14)	0.0049 (15)	0.0023 (14)
C3	0.0107 (18)	0.0303 (19)	0.0229 (18)	-0.0003 (14)	0.0089 (14)	0.0090 (14)
C4	0.0163 (19)	0.0298 (19)	0.0294 (19)	0.0059 (15)	0.0151 (15)	0.0007 (15)
C5	0.0183 (19)	0.0183 (17)	0.0190 (17)	0.0011 (13)	0.0103 (14)	0.0003 (13)
C6	0.0131 (17)	0.0135 (15)	0.0115 (15)	0.0032 (13)	0.0047 (13)	0.0026 (12)
C7	0.0147 (18)	0.0117 (15)	0.0119 (15)	0.0027 (12)	0.0079 (13)	-0.0001 (12)
C8	0.0187 (19)	0.0152 (16)	0.0189 (17)	0.0012 (13)	0.0084 (14)	-0.0001 (13)
C9	0.021 (2)	0.0164 (16)	0.0146 (16)	0.0014 (14)	0.0087 (14)	0.0013 (13)
C10	0.0117 (16)	0.0101 (15)	0.0154 (16)	-0.0049 (12)	0.0090 (14)	0.0009 (13)

*Geometric parameters (Å, °)*

Br1—C3	1.897 (3)	C5—C6	1.391 (4)
O1—C10	1.245 (3)	C5—H5	0.9500
N1—C10	1.332 (3)	C6—C7	1.537 (4)
N1—H1A	0.886 (9)	C7—C9	1.536 (4)
N1—H1B	0.882 (9)	C7—C8	1.540 (4)
C1—C2	1.373 (4)	C7—C10	1.547 (4)
C1—C6	1.396 (4)	C8—H8A	0.9800
C1—H1	0.9500	C8—H8B	0.9800
C2—C3	1.387 (4)	C8—H8C	0.9800
C2—H2	0.9500	C9—H9A	0.9800
C3—C4	1.387 (4)	C9—H9B	0.9800
C4—C5	1.383 (4)	C9—H9C	0.9800
C4—H4	0.9500		
C10—N1—H1A	125.1 (16)	C9—C7—C6	109.3 (2)
C10—N1—H1B	117.5 (15)	C9—C7—C8	108.9 (2)
H1A—N1—H1B	117.2 (16)	C6—C7—C8	112.7 (2)
C2—C1—C6	121.5 (3)	C9—C7—C10	109.2 (2)
C2—C1—H1	119.2	C6—C7—C10	107.4 (2)
C6—C1—H1	119.2	C8—C7—C10	109.3 (2)
C1—C2—C3	119.7 (3)	C7—C8—H8A	109.5
C1—C2—H2	120.1	C7—C8—H8B	109.5
C3—C2—H2	120.1	H8A—C8—H8B	109.5
C4—C3—C2	120.0 (3)	C7—C8—H8C	109.5
C4—C3—Br1	120.3 (2)	H8A—C8—H8C	109.5
C2—C3—Br1	119.6 (2)	H8B—C8—H8C	109.5
C5—C4—C3	119.5 (3)	C7—C9—H9A	109.5
C5—C4—H4	120.3	C7—C9—H9B	109.5
C3—C4—H4	120.3	H9A—C9—H9B	109.5
C4—C5—C6	121.5 (3)	C7—C9—H9C	109.5
C4—C5—H5	119.3	H9A—C9—H9C	109.5
C6—C5—H5	119.3	H9B—C9—H9C	109.5
C5—C6—C1	117.7 (3)	O1—C10—N1	121.1 (3)

C5—C6—C7	122.4 (3)	O1—C10—C7	121.6 (2)
C1—C6—C7	119.9 (3)	N1—C10—C7	117.3 (2)
C6—C1—C2—C3	0.0 (5)	C1—C6—C7—C9	-78.0 (3)
C1—C2—C3—C4	-3.2 (5)	C5—C6—C7—C8	-19.2 (4)
C1—C2—C3—Br1	173.2 (2)	C1—C6—C7—C8	160.7 (2)
C2—C3—C4—C5	3.1 (5)	C5—C6—C7—C10	-139.6 (3)
Br1—C3—C4—C5	-173.2 (2)	C1—C6—C7—C10	40.3 (3)
C3—C4—C5—C6	0.1 (5)	C9—C7—C10—O1	14.6 (4)
C4—C5—C6—C1	-3.1 (4)	C6—C7—C10—O1	-103.8 (3)
C4—C5—C6—C7	176.8 (3)	C8—C7—C10—O1	133.6 (3)
C2—C1—C6—C5	3.0 (4)	C9—C7—C10—N1	-165.7 (2)
C2—C1—C6—C7	-176.8 (3)	C6—C7—C10—N1	75.9 (3)
C5—C6—C7—C9	102.1 (3)	C8—C7—C10—N1	-46.6 (3)

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N1—H1A $\cdots$ O1 <sup>i</sup>	0.89 (1)	2.12 (1)	2.990 (3)	167 (3)
N1—H1B $\cdots$ O1 <sup>ii</sup>	0.88 (1)	2.12 (1)	3.002 (3)	173 (3)

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