

(E)-N'-(5-Bromo-2-methoxybenzylidene)-2-chlorobenzohydrazide

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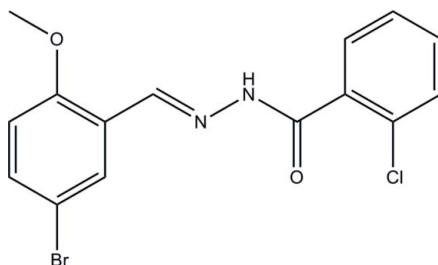
Received 27 July 2011; accepted 23 August 2011

Key indicators: single-crystal X-ray study; $T = 298\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.005\text{ \AA}$;
 R factor = 0.036; wR factor = 0.094; data-to-parameter ratio = 12.6.

In the title compound, $\text{C}_{15}\text{H}_{12}\text{BrClN}_2\text{O}_2$, the dihedral angle between the two substituted aromatic rings is $77.8(3)^\circ$. The molecule exists in a *trans* conformation with respect to the methyldiene unit. In the crystal structure, inversion dimers linked by pairs of $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds generate $R_2^8(8)$ loops.

Related literature

For the crystal structures of some related hydrazone compounds, see: Li (2011a,b); Hashemian *et al.* (2011); Lei (2011); Shalash *et al.* (2010). For hydrogen-bond notation, see: Bernstein *et al.* (1995).



Experimental

Crystal data

$\text{C}_{15}\text{H}_{12}\text{BrClN}_2\text{O}_2$

$M_r = 367.63$

Monoclinic, P_{2_1}/c
 $a = 11.312(2)\text{ \AA}$

$b = 7.374(2)\text{ \AA}$

$c = 17.979(3)\text{ \AA}$

$\beta = 91.972(3)^\circ$

$V = 1499.0(5)\text{ \AA}^3$

$Z = 4$
Mo $K\alpha$ radiation
 $\mu = 2.93\text{ mm}^{-1}$

$T = 298\text{ K}$
 $0.12 \times 0.10 \times 0.07\text{ mm}$

Data collection

Bruker SMART CCD
diffractometer
Absorption correction: multi-scan
(*SADABS*; Sheldrick, 1996)
 $T_{\min} = 0.720$, $T_{\max} = 0.821$

5249 measured reflections
2451 independent reflections
1877 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.050$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.036$
 $wR(F^2) = 0.094$
 $S = 1.04$
2451 reflections
194 parameters
1 restraint

H atoms treated by a mixture of
independent and constrained
refinement
 $\Delta\rho_{\max} = 0.46\text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.46\text{ e \AA}^{-3}$

Table 1
Hydrogen-bond geometry (\AA , $^\circ$).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
N2—H2 \cdots O2 ⁱ	0.89 (1)	1.99 (1)	2.882 (4)	175 (4)

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

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINT* (Bruker, 1998); data reduction: *SAINT*; 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: *SHELXTL*.

The author acknowledges the Zibo Vocational Institute for supporting this work.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB6337).

References

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supporting information

Acta Cryst. (2011). E67, o2511 [doi:10.1107/S1600536811034623]

(E)-N'-(5-Bromo-2-methoxybenzylidene)-2-chlorobenzohydrazide

Xiao-Yan Li

S1. Comment

In the last few years, hydrazones have been attracted much attention for their crystal structures (e.g. Li, 2011*a*; Hashemian *et al.*, 2011; Lei, 2011; Shalash *et al.*, 2010). The author reports herein the crystal structure of the title new hydrazone compound, (I).

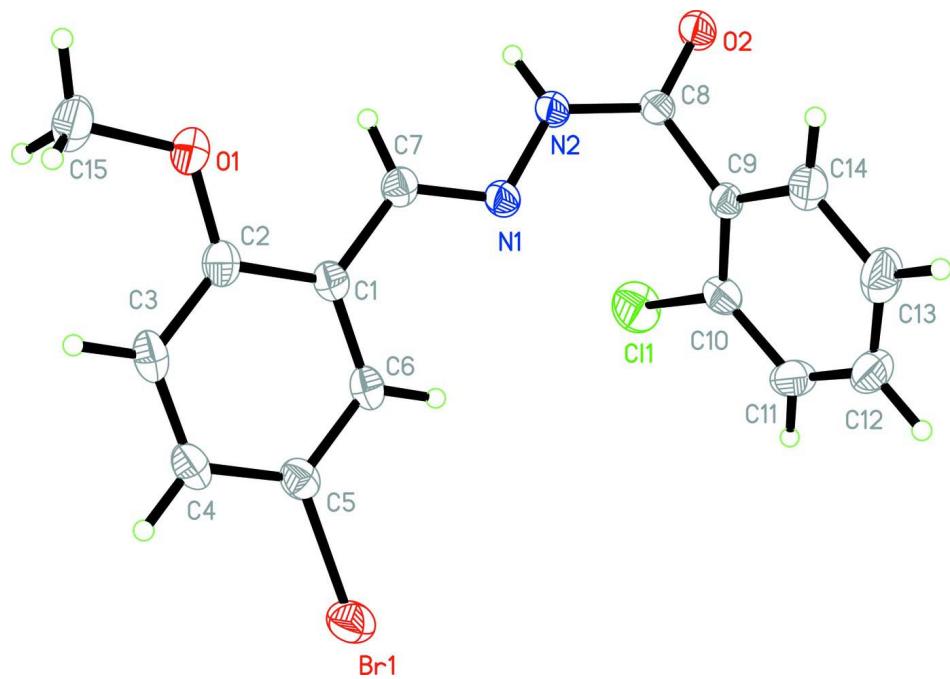
In the title compound, Fig. 1, the dihedral angle between the two substituted aromatic rings is 77.8 (3)°. The molecule exists in a *trans* configuration with respect to the methylidene unit. The bond values are comparable to those observed in a similar compound the author reported recently (Li, 2011*b*). In the crystal structure, adjacent two molecules are linked through two intermolecular N—H···O hydrogen bonds (Table 1), forming a dimer (Fig. 2).

S2. Experimental

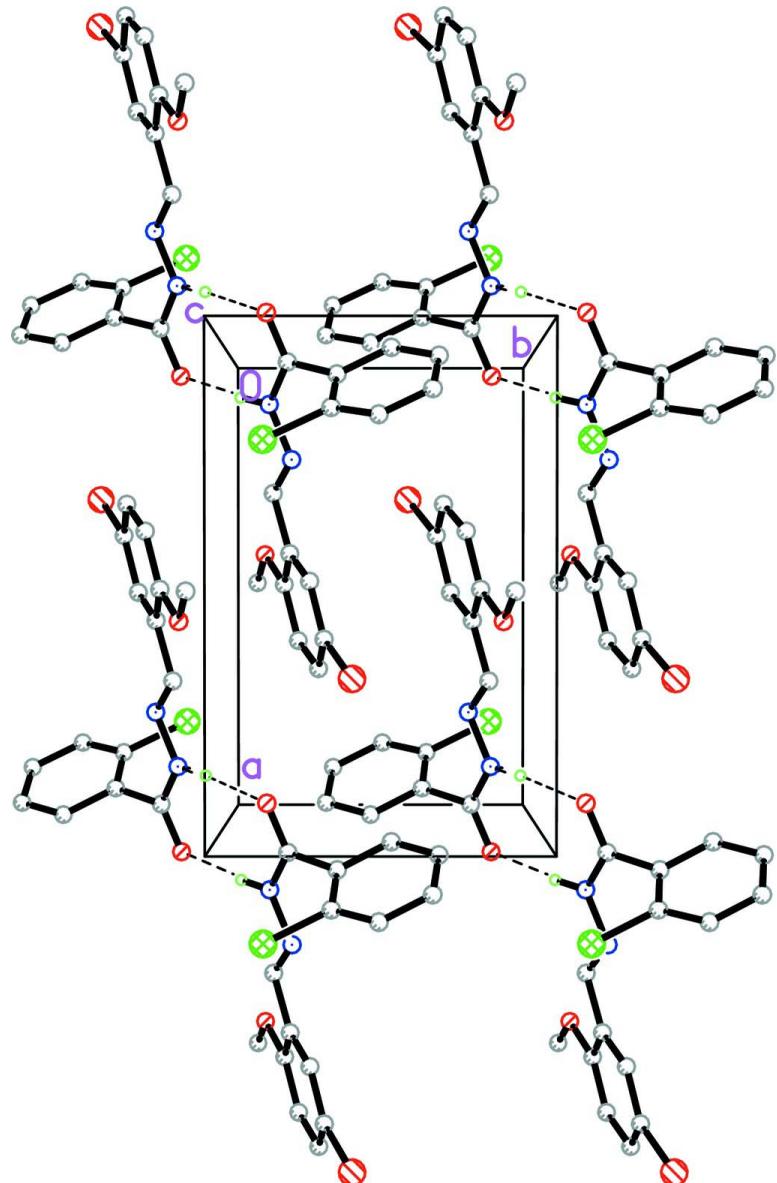
A mixture of 2-chlorobenzhydrazide (0.171 g, 1 mmol) and 5-bromo-2-methoxybenzaldehyde (0.215 g, 1 mmol) in 30 ml of ethanol containing few drops of acetic acid was refluxed for about 1 h. On cooling to room temperature, a solid precipitate was formed. The solid was filtered and then recrystallized from methanol. Colorless blocks of (I) were obtained by slow evaporation of the solution.

S3. Refinement

The N-bound H atom was located from a difference Fourier map and refined isotropically. The rest of H atoms were positioned geometrically [C—H = 0.93 and 0.96 Å] and refined using a riding model [$U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ and $1.5U_{\text{eq}}(\text{C}15)$]. A rotating-group model was applied for the methyl group.

**Figure 1**

The molecular structure of (I) at 30% probability level. Hydrogen bonds are indicated by dashed lines.

**Figure 2**

The packing of (I), viewed along the c axis. Intermolecular hydrogen bonds are drawn as dashed lines.

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 Monoclinic, $P2_1/c$
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 $c = 17.979 (3) \text{ \AA}$
 $\beta = 91.972 (3)^\circ$
 $V = 1499.0 (5) \text{ \AA}^3$
 $Z = 4$

$F(000) = 736$
 $D_x = 1.629 \text{ Mg m}^{-3}$
 Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$
 Cell parameters from 1972 reflections
 $\theta = 3.1\text{--}24.9^\circ$
 $\mu = 2.93 \text{ mm}^{-1}$
 $T = 298 \text{ K}$
 Block, colourless
 $0.12 \times 0.10 \times 0.07 \text{ mm}$

Data collection

Bruker SMART CCD
diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
 ω scans
Absorption correction: multi-scan
(SADABS; Sheldrick, 1996)
 $T_{\min} = 0.720$, $T_{\max} = 0.821$

5249 measured reflections
2451 independent reflections
1877 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.050$
 $\theta_{\max} = 25.0^\circ$, $\theta_{\min} = 2.9^\circ$
 $h = -13 \rightarrow 13$
 $k = -8 \rightarrow 6$
 $l = -20 \rightarrow 18$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.036$
 $wR(F^2) = 0.094$
 $S = 1.04$
2451 reflections
194 parameters
1 restraint
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
 $w = 1/[\sigma^2(F_o^2) + (0.0429P)^2 + 0.678P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.46 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.46 \text{ e } \text{\AA}^{-3}$

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 $F^2 > 2\text{sigma}(F^2)$ 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>	$U_{\text{iso}}^*/U_{\text{eq}}$
Br1	0.68536 (3)	0.08870 (6)	0.13010 (2)	0.05792 (18)
Cl1	0.20872 (10)	0.35802 (17)	0.20909 (7)	0.0738 (4)
N1	0.2386 (2)	0.2805 (4)	0.03030 (15)	0.0362 (6)
N2	0.1252 (2)	0.3513 (4)	0.02797 (16)	0.0405 (7)
O1	0.4312 (2)	0.3783 (4)	-0.14532 (14)	0.0534 (7)
O2	-0.05797 (18)	0.3567 (3)	0.07125 (14)	0.0533 (7)
C1	0.4308 (2)	0.2898 (4)	-0.02015 (18)	0.0358 (7)
C2	0.4949 (3)	0.3117 (5)	-0.08532 (19)	0.0402 (8)
C3	0.6132 (3)	0.2662 (5)	-0.0855 (2)	0.0484 (9)
H3	0.6552	0.2811	-0.1287	0.058*
C4	0.6700 (3)	0.1984 (5)	-0.0217 (2)	0.0468 (9)
H4	0.7496	0.1668	-0.0221	0.056*
C5	0.6074 (3)	0.1783 (5)	0.04254 (19)	0.0401 (8)
C6	0.4885 (3)	0.2236 (4)	0.04331 (19)	0.0390 (8)
H6	0.4472	0.2093	0.0868	0.047*

C7	0.3058 (3)	0.3422 (4)	-0.01972 (19)	0.0375 (8)
H7	0.2752	0.4208	-0.0560	0.045*
C8	0.0425 (2)	0.2909 (4)	0.07350 (18)	0.0362 (7)
C9	0.0717 (2)	0.1349 (4)	0.12478 (18)	0.0348 (8)
C10	0.1435 (3)	0.1509 (5)	0.1879 (2)	0.0445 (9)
C11	0.1611 (3)	0.0055 (7)	0.2367 (2)	0.0578 (11)
H11	0.2078	0.0194	0.2799	0.069*
C12	0.1087 (4)	-0.1580 (6)	0.2202 (3)	0.0627 (12)
H12	0.1215	-0.2560	0.2521	0.075*
C13	0.0377 (4)	-0.1789 (6)	0.1573 (3)	0.0613 (11)
H13	0.0037	-0.2910	0.1465	0.074*
C14	0.0166 (3)	-0.0332 (5)	0.1098 (2)	0.0476 (9)
H14	-0.0338	-0.0464	0.0682	0.057*
C15	0.4912 (4)	0.4085 (6)	-0.2124 (2)	0.0649 (12)
H15A	0.5509	0.4995	-0.2042	0.097*
H15B	0.4357	0.4488	-0.2505	0.097*
H15C	0.5276	0.2977	-0.2278	0.097*
H2	0.109 (4)	0.444 (4)	-0.0024 (19)	0.080*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Br1	0.0437 (2)	0.0720 (3)	0.0575 (3)	0.00185 (18)	-0.00739 (17)	0.0012 (2)
Cl1	0.0748 (7)	0.0728 (8)	0.0728 (8)	-0.0254 (6)	-0.0124 (6)	-0.0101 (6)
N1	0.0305 (12)	0.0354 (16)	0.0430 (17)	0.0034 (11)	0.0047 (11)	0.0041 (13)
N2	0.0293 (13)	0.0413 (17)	0.051 (2)	0.0057 (11)	0.0055 (12)	0.0133 (14)
O1	0.0489 (14)	0.0648 (18)	0.0473 (16)	0.0000 (12)	0.0109 (12)	0.0140 (13)
O2	0.0318 (12)	0.0558 (16)	0.0728 (19)	0.0101 (11)	0.0087 (11)	0.0244 (14)
C1	0.0330 (15)	0.0330 (18)	0.042 (2)	-0.0017 (13)	0.0054 (14)	-0.0016 (15)
C2	0.0435 (18)	0.035 (2)	0.042 (2)	-0.0066 (15)	0.0062 (15)	-0.0031 (16)
C3	0.0410 (18)	0.051 (2)	0.054 (2)	-0.0043 (16)	0.0197 (17)	-0.0017 (19)
C4	0.0323 (16)	0.048 (2)	0.061 (3)	-0.0004 (15)	0.0104 (16)	-0.0026 (19)
C5	0.0356 (16)	0.0345 (19)	0.050 (2)	-0.0040 (14)	0.0011 (15)	-0.0035 (16)
C6	0.0347 (16)	0.0371 (19)	0.046 (2)	-0.0060 (14)	0.0086 (15)	-0.0037 (16)
C7	0.0392 (17)	0.0310 (18)	0.042 (2)	0.0000 (13)	0.0045 (15)	0.0010 (15)
C8	0.0312 (15)	0.0343 (19)	0.043 (2)	-0.0012 (13)	-0.0012 (13)	0.0042 (15)
C9	0.0285 (15)	0.0348 (19)	0.042 (2)	0.0046 (12)	0.0089 (14)	0.0051 (15)
C10	0.0339 (16)	0.050 (2)	0.049 (2)	0.0010 (15)	0.0013 (16)	-0.0007 (18)
C11	0.047 (2)	0.075 (3)	0.051 (3)	0.014 (2)	0.0002 (17)	0.017 (2)
C12	0.056 (2)	0.062 (3)	0.071 (3)	0.012 (2)	0.012 (2)	0.031 (2)
C13	0.067 (3)	0.039 (2)	0.079 (3)	-0.0035 (19)	0.017 (2)	0.010 (2)
C14	0.0476 (19)	0.044 (2)	0.051 (2)	-0.0011 (16)	0.0098 (17)	0.0049 (18)
C15	0.066 (2)	0.079 (3)	0.050 (3)	-0.011 (2)	0.014 (2)	0.012 (2)

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

Br1—C5	1.897 (3)	C5—C6	1.386 (4)
Cl1—C10	1.733 (4)	C6—H6	0.9300

N1—C7	1.282 (4)	C7—H7	0.9300
N1—N2	1.384 (3)	C8—C9	1.504 (4)
N2—C8	1.340 (4)	C9—C10	1.377 (5)
N2—H2	0.893 (10)	C9—C14	1.409 (5)
O1—C2	1.368 (4)	C10—C11	1.395 (5)
O1—C15	1.422 (4)	C11—C12	1.371 (6)
O2—C8	1.235 (3)	C11—H11	0.9300
C1—C6	1.384 (5)	C12—C13	1.373 (6)
C1—C2	1.408 (4)	C12—H12	0.9300
C1—C7	1.466 (4)	C13—C14	1.388 (5)
C2—C3	1.380 (5)	C13—H13	0.9300
C3—C4	1.390 (5)	C14—H14	0.9300
C3—H3	0.9300	C15—H15A	0.9600
C4—C5	1.384 (5)	C15—H15B	0.9600
C4—H4	0.9300	C15—H15C	0.9600
C7—N1—N2	114.6 (3)	O2—C8—C9	120.2 (3)
C8—N2—N1	121.5 (3)	N2—C8—C9	119.0 (3)
C8—N2—H2	120 (3)	C10—C9—C14	118.5 (3)
N1—N2—H2	119 (3)	C10—C9—C8	123.4 (3)
C2—O1—C15	118.1 (3)	C14—C9—C8	118.0 (3)
C6—C1—C2	119.0 (3)	C9—C10—C11	121.3 (4)
C6—C1—C7	121.1 (3)	C9—C10—Cl1	119.4 (3)
C2—C1—C7	119.9 (3)	C11—C10—Cl1	119.2 (3)
O1—C2—C3	124.7 (3)	C12—C11—C10	119.3 (4)
O1—C2—C1	115.2 (3)	C12—C11—H11	120.4
C3—C2—C1	120.1 (3)	C12—C11—H11	120.4
C2—C3—C4	120.4 (3)	C11—C12—C13	120.8 (4)
C2—C3—H3	119.8	C11—C12—H12	119.6
C4—C3—H3	119.8	C13—C12—H12	119.6
C5—C4—C3	119.5 (3)	C12—C13—C14	120.2 (4)
C5—C4—H4	120.2	C12—C13—H13	119.9
C3—C4—H4	120.2	C14—C13—H13	119.9
C4—C5—C6	120.5 (3)	C13—C14—C9	119.9 (4)
C4—C5—Br1	119.5 (2)	C13—C14—H14	120.1
C6—C5—Br1	120.0 (3)	C9—C14—H14	120.1
C1—C6—C5	120.5 (3)	O1—C15—H15A	109.5
C1—C6—H6	119.8	O1—C15—H15B	109.5
C5—C6—H6	119.8	H15A—C15—H15B	109.5
N1—C7—C1	120.4 (3)	O1—C15—H15C	109.5
N1—C7—H7	119.8	H15A—C15—H15C	109.5
C1—C7—H7	119.8	H15B—C15—H15C	109.5
O2—C8—N2	120.7 (3)		

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H···A
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supporting information

N2—H2···O2 ⁱ	0.89 (1)	1.99 (1)	2.882 (4)	175 (4)
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Symmetry code: (i) $-x, -y+1, -z$.