



Synthesis, X-ray crystal structure, Hirshfeld surface analysis and DFT studies of (*E*)-*N'*-(2-bromobenzylidene)-4-methylbenzohydrazide

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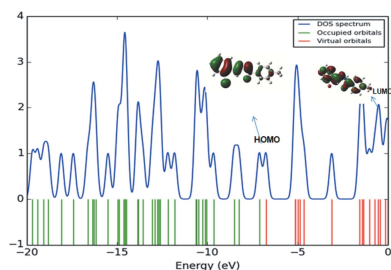
Keywords: crystal structure; hydrogen bonding; DFT; Hirshfeld surface analysis.**CCDC reference:** 1580647**Supporting information:** this article has supporting information at journals.iucr.org/e

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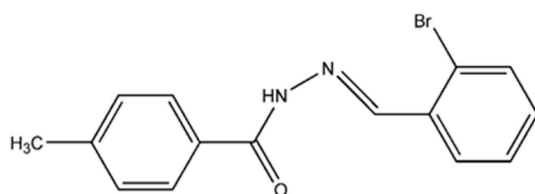
The title molecule, C₁₅H₁₃BrN₂O, displays a *trans* configuration with respect to the C=N double bond. The dihedral angle between the bromo- and methyl-substituted benzene rings is 16.1 (3)°. In the crystal, molecules are connected by N—H···O and weak C—H···O hydrogen bonds, forming R₂¹(6) ring motifs and generating chains along the *a*-axis direction. The optimized structure generated theoretically *via* density functional theory (DFT) using standard B3LYP functional and 6–311 G(d,p) basis-set calculations renders good support to the experimental data. The HOMO–LUMO behaviour was elucidated to determine the energy gap. The intermolecular interactions were quantified and analysed using Hirshfeld surface analysis.

1. Chemical context

Hydrazones are a class of organic compounds that possess an R₁R₂C=NNH₂ structural motif. They are related to ketones and aldehydes in which oxygen has been replaced with an NNH₂ group (Rollas & Küçükgül, 2007). Azomethines, –NHN=CH–, constitute an important class of compounds for new drug development. The reaction of a hydrazine or hydrazide with aldehydes and ketones yields hydrazones. Hydrazones are important in drug design as they act as ligands for metal complexes, organocatalysis and the synthesis of organic compounds. The C=N bond of the hydrazone and the terminal nitrogen atom containing a lone pair of electron is responsible for the physical and chemical properties. The C atom in the hydrazone unit has both electrophilic and nucleophilic character and both the N atoms are nucleophilic, although the amino-type nitrogen is more reactive. As a result of these properties, hydrazones are widely used in organic synthesis. Owing to their ease of preparation and diverse pharmacological potential, much work on hydrazones has been carried out by medicinal chemists to develop agents with better activity and low toxicity profiles. Hydrazones are known to possess diverse biological activities such as antimicrobial, anti-inflammatory, anticancer and antimalarial (Yousef *et al.*, 2003; Trepanier *et al.*, 1966) and have been evaluated for inhibition of PDE10A, a phosphodiesterase responsible for neurobiological and psychological disorders such as Parkinson's, schizophrenia and Huntington's disease (Gage *et al.*, 2011). The anticonvulsant potential of some hydrazone derivatives having long duration and rapid onset of action have been reported (Kaushik *et al.*, 2010), as has their anti-depressant activity (de Oliveira *et al.*, 2011).



Schiff bases are used widely in the field of coordination chemistry and have interesting properties (Morshedi *et al.*, 2009; Zhou *et al.*, 2006; Khanmohammadi *et al.*, 2009). These compounds are synthesized by condensation of carbonyl compounds with amines (van den Ancker *et al.*, 2006; Hamaker *et al.*, 2010). In addition, free Schiff base compounds are reported to possess antimicrobial (Aslantas *et al.*, 2009) and non-linear optical (Karakas *et al.*, 2008) properties. Our previous work on (*E*)-4-bromo-*N'*-(2,4-dihydroxybenzylidene)benzohydrazide and (*E*)-4-toluic-*N'*-(2,4-dihydroxybenzylidene)benzohydrazide have been recently reported (Arunagiri *et al.*, 2018*a,b*). This work has been a guide for the development of the new Schiff base title compound, which possesses electronic and non-linear properties. As part of our interest in the identification of bioactive compounds, we report herein on its crystal structure.



2. Structural commentary

The molecular structure of the title compound is shown in Fig. 1(*a*). The molecule adopts an (*E*) configuration across the C=N bond, joining the hydrazide group and the benzene ring. In the crystal, the dihedral angle between the bromo- and methyl-substituted benzene rings is 16.1 (3)°. The structure was optimized with the *Gaussian09W* software (Frisch *et al.*, 2009) using the DFT-B3LYP/6-311G(d,p) method, providing

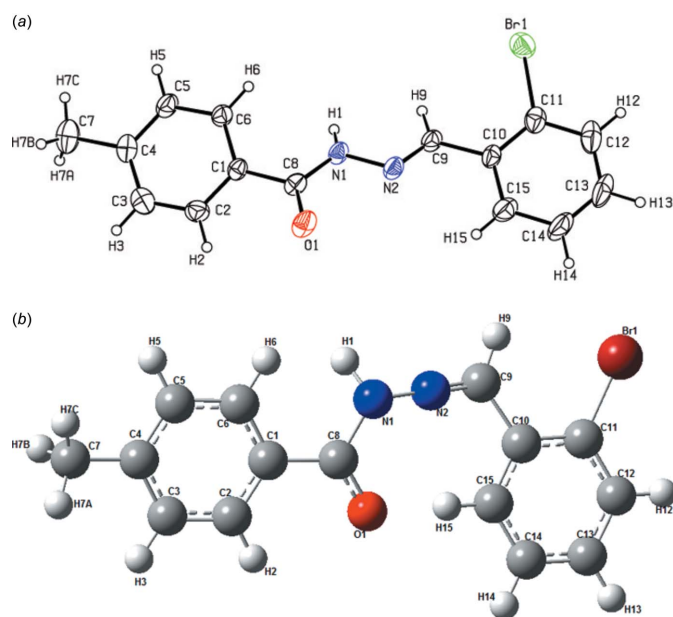


Figure 1
(*a*) The molecular structure of the title compound, with displacement ellipsoids drawn at the 50% probability level. (*b*) The optimized structure of the title compound.

Table 1
Selected geometric parameters (Å, °) for the experimental and DFT structures.

	XRD	DFT
Br1—C11	1.891 (6)	1.925
O1—C8	1.225 (6)	1.219
N1—N2	1.379 (5)	1.364
N1—C8	1.351 (6)	1.383
N2—C9	1.270 (6)	1.285
N1—H1	0.837 (19)	1.006
C1—C8	1.485 (6)	1.501
C9—C10	1.470 (8)	1.470
C9—H9	0.930	1.082
N2—N1—C8	119.2 (4)	120.00
C8—N1—H1	123 (4)	117.84
N2—N1—H1	117 (4)	111.31
N1—C8—C1	116.6 (4)	114.46
O1—C8—C1	121.6 (4)	122.20
O1—C8—N1	121.7 (4)	123.33
N2—C9—C10	120.1 (4)	118.62
N2—C9—H9	120.00	122.79
C8—N1—N2—C9	−173.3 (5)	−179.68
N2—N1—C8—O1	−3.5 (7)	2.54
N2—N1—C8—C1	174.64	−178.45
C2—C1—C8—O1	24.6 (7)	22.94
C2—C1—C8—N1	−153.6 (5)	−156.06
N2—C9—C10—C11	161.4 (5)	−179.93
N2—C9—C10—C15	−19.2 (7)	−0.14

Table 2
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>
N1—H1...O1 ⁱ	0.84 (3)	1.96 (3)	2.785 (5)	167 (5)
C9—H9...O1 ⁱ	0.93	2.38	3.166 (6)	142

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

information about the geometry of the molecule. The optimized structure is shown in Fig. 1(*b*). The geometrical parameters (Table 1) are mostly within normal ranges, the slight deviations of the theoretical values from those determined experimentally are due to the fact that the optimization is performed in isolated conditions, whereas the crystal environment and hydrogen-bonding interactions affect the results of the X-ray structure (Zainuri *et al.*, 2017).

The hydrazide unit (N1/N2/C1/C8—C10) is essentially planar, with a maximum deviation from the least-squares plane of 0.099 (4) Å for atom C10. The O1=C8 bond length [1.225 (6) and 1.219 Å for XRD and B3LYP, respectively] indicates single-bond character. The N1—N2 bond length [1.379 (5) Å for XRD and 1.364 Å for B3LYP] is in good agreement with other experimental values (Sivajeyanthi *et al.*, 2017). The C—N bond lengths range from a typical single bond [C8—N1 = 1.351 (6) Å] to a double bond [C9=N2 = 1.270 (6) Å (Sivajeyanthi *et al.*, 2017; Arunagiri *et al.*, 2018*a,b*).

3. Supramolecular features

In the crystal, N1—H1...O1ⁱ and C9—H9...O1ⁱ hydrogen bonds (Table 2) connect symmetry-related molecules through

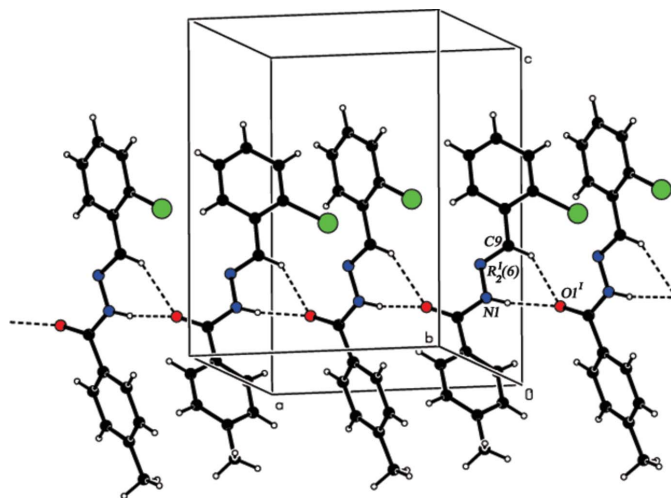


Figure 2
Part of the crystal structure with hydrogen bonds shown as dashed lines.

classical N—H···O and weak C—H···O hydrogen bonds, forming $R_2^1(6)$ ring motifs and generating [100] chains (Fig. 2).

4. Hirshfeld surface analysis

Hirshfeld surface analysis (McKinnon *et al.*, 2007; Spackman & Jayatilaka, 2009) along with decomposed 2D fingerprint plots (Spackman & McKinnon, 2002; McKinnon *et al.*, 2004, 2007) mapped over d_{norm} , shape-index and curvedness were used to visualize and quantify the intermolecular interactions. The Hirshfeld surface (HS) and fingerprint plots were generated based on the d_i and d_e distances using *Crystal Explorer3.1* (Wolff *et al.*, 2012) where d_i is the distance from the nearest atom inside the surface, while d_e is the distance from the HS to the nearest atom outside the surface. In the d_{norm} surfaces, large circular depressions (deep red) are the indicators of hydrogen-bonding contacts whereas other visible spots are due to H···H contacts. The dominant H···O interaction in the title compound is evident as a bright-red area in Fig. 3 while the light-red spots are due to N—H···O and C—H···O interactions. The shape-index surface [Fig. 4(a)] conveys information about each donor–acceptor pair and while the

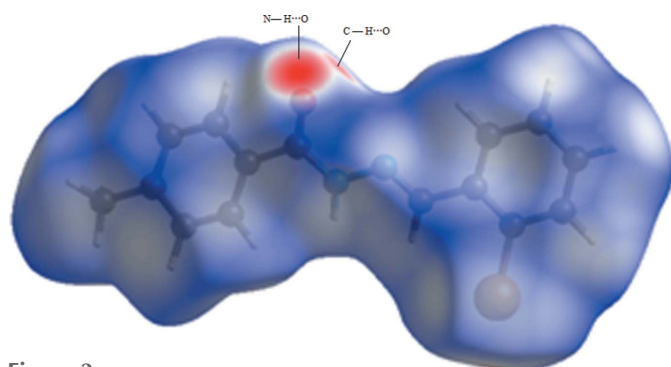


Figure 3
The three-dimensional d_{norm} surface of the title compound. **add contouring levels**

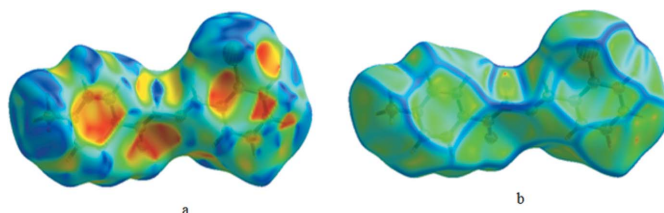


Figure 4
Hirshfeld surfaces mapped over (a) shape-index and (b) curvedness for the title compound.

curvedness surface [Fig. 4(b)] is effectively divided into sets of patches, respectively. The tiny extent of area and light colour on the surface indicates weaker and longer contacts other than hydrogen bonds. The 2D fingerprint plots in Fig. 5 shows the relative contributions from the various intermolecular contacts (O···H, H···H, C···H, C···C, N···H, N···N, O···Br and C···Br) in the crystal structure. The H···H contacts (36%) make the largest contribution, followed by C···H/H···C (28.2%), O···H/H···O (10.2%) and N···H/H···N (7.5%), the latter interactions being represented by blue spikes on both sides at the bottom of the plot.

5. Frontier molecular orbitals and Molecular electrostatic potential analysis

The highest-occupied molecular orbital (HOMO), which acts as an electron donor, and the lowest-unoccupied molecular orbital (LUMO), which acts as an electron acceptor, are very important parameters for quantum chemistry. If the energy gap is small, then the molecule is highly polarizable and has

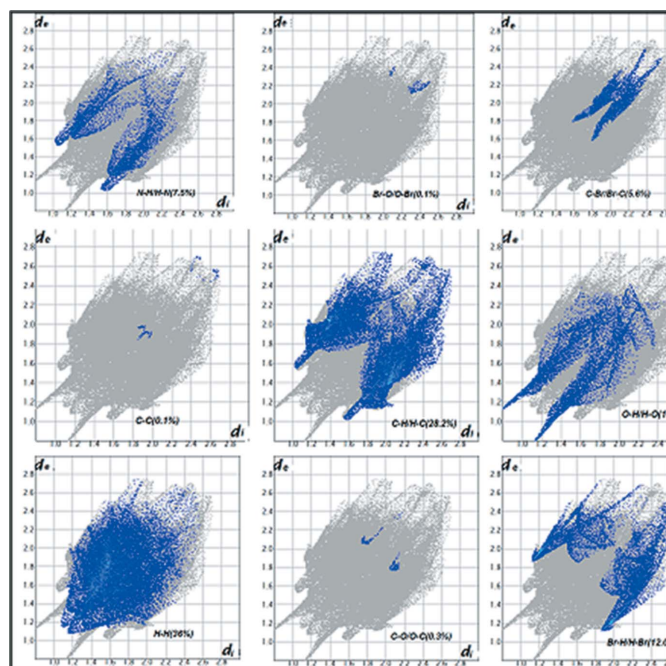


Figure 5
Two-dimensional fingerprint plots with the relative contributions of the various interactions.

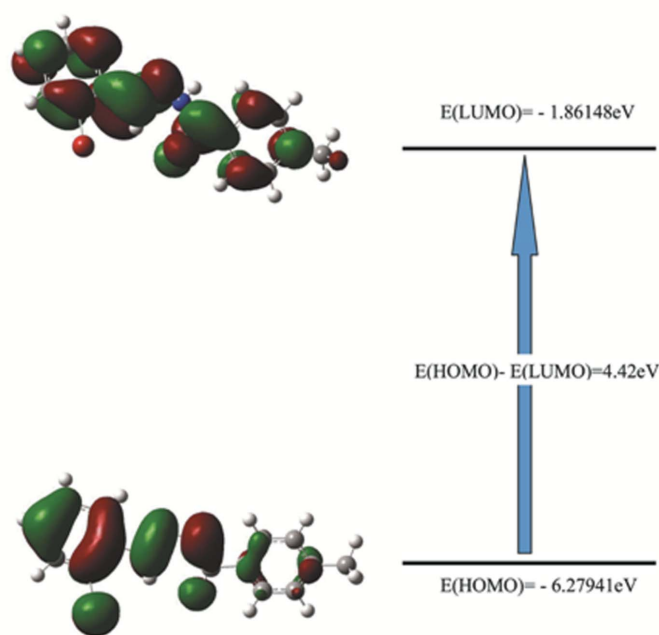


Figure 6
The energy band gap of the compound.

high chemical reactivity. The energy levels were computed by the DFTB3LYP/6-311G(d,p) method (Becke *et al.*, 1993) as implemented in *GAUSSIAN09W* (Frisch *et al.*, 2009). The electron transition from the HOMO to the LUMO energy level is shown in Fig. 6. The molecular orbital of HOMO contain both σ and π electron-density character, whereas the LUMO is mainly composed of π -orbital density. The energy band gap (ΔE) of the molecule is about 4.42 eV.

The *Gauss-Sum2.2* program (O'Boyle *et al.*, 2008) was used to calculate group contributions to the molecular orbitals (HOMO and LUMO) and prepare the density of states (DOS) spectrum shown in Fig. 7. The DOS spectrum was formed by convoluting the molecular orbital information with GAUSSIAN curves of unit height. The green and red lines in the DOS spectrum indicate the HOMO and LUMO levels. The

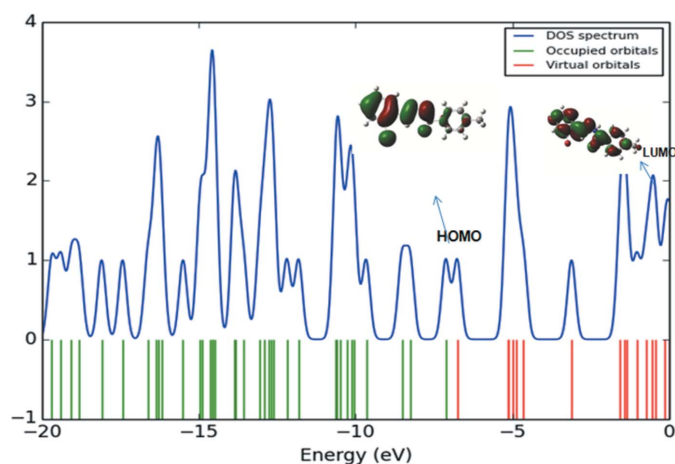


Figure 7
The density of states (DOS) spectrum of the compound.

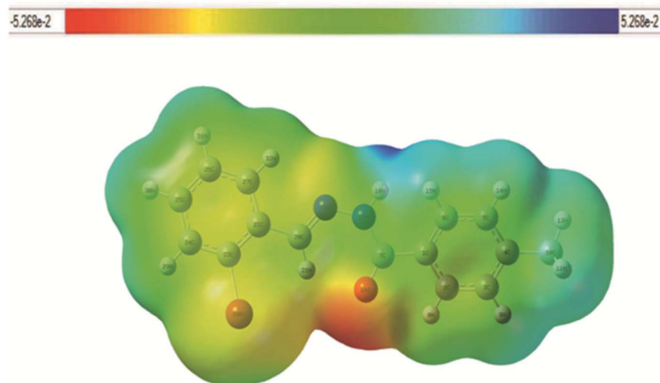


Figure 8
The molecular electrostatic potential map for the title compound.

DOS spectrum supports the energy gap calculated by HOMO–LUMO analysis. A molecule with a large energy gap is described as hard while one having a small energy gap is known as a soft molecule. Hard molecules are not more polarizable than the soft ones because they require immense excitation energy (Karabacak & Yilan, 2012).

The molecular electrostatic potential is related to the electron density and molecular electrostatic potential (MESP) maps are very useful descriptors for understanding reactive sites for electrophilic and nucleophilic reactions as well as hydrogen-bonding interactions (Sebastian & Sundaraganesan, 2010; Luque *et al.*, 2000). Different values of the electrostatic potential are represented by different colours: red represents regions of the most electronegative electrostatic potential, blue represents regions of the most positive electrostatic potential and green represents regions of zero potential. The potential increases in the following order: red < orange < yellow < green < blue. Herein, MEP was calculated at the DFT–B3LYP/6–311(d,p) level of theory that was used for optimization. The MESP map for the title molecule is shown in Fig. 8 with a colour range from -0.053 (red) to 0.053 a.u. (blue). The most electrostatically positive region (blue colour) is located in the molecular plane (N-bonded hydrogen atoms of toluic hydrazide), thus explaining N1–H1...O1¹ hydrogen bond observed in the crystal structure. The map clearly shows that the electron-rich (red) region is spread around the carbonyl oxygen atom whereas the hydrogen atom attached to nitrogen is positively charged (blue).

6. Database survey

A search of the Cambridge Structural Database (Version 5.39, last update November 2018; Groom *et al.*, 2016) revealed closely related compounds that differ in the donor substituents: *N'*-(4-chlorobenzylidene)-2-hydroxybenzohydrazide (Zhang *et al.*, 2009), (*E*)-*N'*-[(pyridin-2-yl)methylene]benzohydrazide (Ramesh Babu *et al.*, 2014), (*E*)-*N'*-(4-methoxybenzylidene)pyridine-3-carbohydrazide dihydrate (Govindarasu *et al.*, 2015), (*E*)-4-bromo-*N'*-(4-methoxybenzylidene)-benzohydrazide (Balasubramani *et al.*, 2018), (*E*)-3-(1*H*-indol-

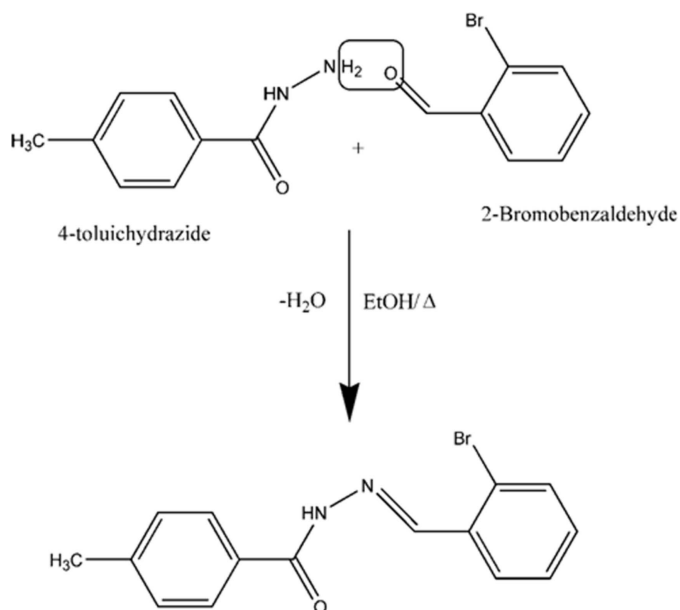


Figure 9
Reaction scheme.

2-yl)-1-(4-nitrophenyl)prop-2-en-1-one hemihydrate (Zaini *et al.*, 2018); (*E*)-4-bromo-*N'*-(2,4-dihydroxybenzylidene)benzohydrazide and (*E*)-4-toluic-*N'*-(2,4-dihydroxybenzylidene)benzohydrazide (Arunagiri *et al.*, 2018*a,b*). In our studies of analogous hydrazide derivatives (with Cl or Br replacing the methyl group of the title compound, we have observed similar types of Intramolecular *S*(6) and intermolecular N—H...O hydrogen bonds (between the amide hydrogen and the carbonyl oxygen atoms).

7. Synthesis and crystallization

The title compound was synthesized by the condensation of 4-toluic hydrazide and 2-bromobenzaldehyde (Fig. 9). An ethanol solution (10 ml) of 4-toluic hydrazide (0.25 mol) was mixed with ethanol solution of 2-bromobenzaldehyde (10 ml, 0.25 mol) and the reaction mixture was heated at 323 K for half an hour with constant stirring before being filtered and kept for crystallization. After a period of one week, brown block-shaped crystals of the title compound were obtained.

8. X-ray crystallography and refinement

Crystal data, data collection and structure refinement details are summarized in Table 3. The hydrogen atom on N1 (H1) was located in a difference-Fourier map and freely refined. C-bound hydrogen atoms were placed in calculated positions (C—H = 0.93–0.96 Å) and refined as riding with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ or $1.5U_{\text{eq}}(\text{C-methyl})$.

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Table 3
Experimental details.

Crystal data	
Chemical formula	$\text{C}_{15}\text{H}_{13}\text{BrN}_2\text{O}$
M_r	317.18
Crystal system, space group	Orthorhombic, $Pna2_1$
Temperature (K)	296
a, b, c (Å)	9.6002 (10), 11.5584 (13), 12.5823 (12)
V (Å ³)	1396.2 (3)
Z	4
Radiation type	Mo $K\alpha$
μ (mm ⁻¹)	2.94
Crystal size (mm)	0.30 × 0.25 × 0.20
Data collection	
Diffractometer	Bruker Kappa APEXII CCD
Absorption correction	Multi-scan (SADABS; Bruker, 2004)
$T_{\text{min}}, T_{\text{max}}$	0.473, 0.591
No. of measured, independent and observed [$I > 2\sigma(I)$] reflections	17816, 2744, 1837
R_{int}	0.035
$(\sin \theta/\lambda)_{\text{max}}$ (Å ⁻¹)	0.617
Refinement	
$R[F^2 > 2\sigma(F^2)], wR(F^2), S$	0.037, 0.097, 1.10
No. of reflections	2744
No. of parameters	176
No. of restraints	2
H-atom treatment	H atoms treated by a mixture of independent and constrained refinement
$\Delta\rho_{\text{max}}, \Delta\rho_{\text{min}}$ (e Å ⁻³)	0.42, -0.47

Computer programs: APEX2, SAINT and XPREP (Bruker, 2004), SIR97 (Altomare *et al.*, 1999), SHELXL014 (Sheldrick, 2015) and ORTEP-3 for Windows (Farrugia, 2012).

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supporting information

Acta Cryst. (2019). E75, 109-114 [https://doi.org/10.1107/S2056989018017978]

Synthesis, X-ray crystal structure, Hirshfeld surface analysis and DFT studies of (E)-N'-(2-bromobenzylidene)-4-methylbenzohydrazide

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Computing details

Data collection: *APEX2* (Bruker, 2004); cell refinement: *APEX2* and *SAINT* (Bruker, 2004); data reduction: *SAINT* and *XPREP* (Bruker, 2004); program(s) used to solve structure: *SIR97* (Altomare *et al.*, 1999); program(s) used to refine structure: *SHELXL014* (Sheldrick, 2015); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 2012); software used to prepare material for publication: *SHELXL2014* (Sheldrick, 2015).

(E)-4-Methyl-N'-(2-bromobenzylidene)benzohydrazide

Crystal data

C₁₅H₁₃BrN₂O

M_r = 317.18

Orthorhombic, *Pna*2₁

Hall symbol: P 2c -2n

a = 9.6002 (10) Å

b = 11.5584 (13) Å

c = 12.5823 (12) Å

V = 1396.2 (3) Å³

Z = 4

F(000) = 640

D_x = 1.509 Mg m⁻³

Mo *Kα* radiation, λ = 0.71073 Å

Cell parameters from 6594 reflections

θ = 2.4–26.0°

μ = 2.94 mm⁻¹

T = 296 K

Block, brown

0.30 × 0.25 × 0.20 mm

Data collection

Bruker Kappa APEXII CCD
diffractometer

Radiation source: fine-focus sealed tube
Graphite monochromator

ω and φ scan

Absorption correction: multi-scan
(SADABS; Bruker, 2004)

T_{min} = 0.473, *T_{max}* = 0.591

17816 measured reflections

2744 independent reflections

1837 reflections with *I* > 2σ(*I*)

R_{int} = 0.035

θ_{max} = 26.0°, θ_{min} = 2.4°

h = -11→11

k = -14→14

l = -15→15

Refinement

Refinement on *F*²

Least-squares matrix: full

R[*F*² > 2σ(*F*²)] = 0.037

wR(*F*²) = 0.097

S = 1.10

2744 reflections

176 parameters

2 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

$$w = 1/[\sigma^2(F_o^2) + (0.0208P)^2 + 2.2419P]$$

where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$

$$\Delta\rho_{\max} = 0.42 \text{ e } \text{\AA}^{-3}$$

$$\Delta\rho_{\min} = -0.47 \text{ e } \text{\AA}^{-3}$$

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)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
Br1	0.27412 (5)	0.49846 (5)	0.50267 (12)	0.06555 (18)
O1	0.7807 (3)	0.1961 (4)	0.2048 (3)	0.0633 (11)
N1	0.5643 (4)	0.2641 (4)	0.2337 (3)	0.0430 (10)
N2	0.5965 (5)	0.2925 (4)	0.3373 (3)	0.0464 (11)
C1	0.6174 (5)	0.1750 (4)	0.0649 (3)	0.0390 (10)
C2	0.6886 (5)	0.0846 (4)	0.0191 (5)	0.0529 (14)
H2	0.7632	0.0514	0.0551	0.064*
C3	0.6512 (7)	0.0427 (5)	-0.0791 (5)	0.0632 (15)
H3	0.7000	-0.0193	-0.1080	0.076*
C4	0.5433 (6)	0.0907 (5)	-0.1352 (4)	0.0546 (14)
C5	0.4721 (6)	0.1827 (5)	-0.0902 (4)	0.0526 (13)
H5	0.3987	0.2165	-0.1271	0.063*
C6	0.5085 (4)	0.2251 (4)	0.0089 (5)	0.0446 (10)
H6	0.4599	0.2872	0.0378	0.054*
C7	0.4998 (11)	0.0447 (6)	-0.2424 (6)	0.0834 (19)
H7C	0.4230	0.0894	-0.2689	0.125*
H7A	0.4723	-0.0348	-0.2356	0.125*
H7B	0.5766	0.0503	-0.2910	0.125*
C8	0.6621 (5)	0.2130 (4)	0.1724 (4)	0.0429 (11)
C9	0.4964 (5)	0.3296 (4)	0.3937 (4)	0.0446 (12)
H9	0.4093	0.3413	0.3632	0.053*
C10	0.5182 (5)	0.3543 (3)	0.5071 (5)	0.0435 (10)
C11	0.4290 (6)	0.4236 (4)	0.5651 (4)	0.0530 (13)
C12	0.4514 (8)	0.4419 (5)	0.6726 (4)	0.0726 (19)
H12	0.3908	0.4882	0.7115	0.087*
C13	0.5642 (9)	0.3908 (6)	0.7213 (5)	0.082 (2)
H13	0.5807	0.4039	0.7931	0.098*
C14	0.6504 (9)	0.3224 (6)	0.6660 (5)	0.083 (2)
H14	0.7239	0.2857	0.7004	0.100*
C15	0.6313 (7)	0.3059 (5)	0.5591 (4)	0.0608 (15)
H15	0.6951	0.2617	0.5210	0.073*
H1	0.480 (3)	0.268 (5)	0.217 (4)	0.051 (16)*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Br1	0.0673 (3)	0.0624 (3)	0.0669 (3)	0.0056 (3)	0.0118 (6)	-0.0006 (3)
O1	0.0301 (18)	0.111 (3)	0.049 (2)	0.006 (2)	-0.0069 (16)	0.001 (2)
N1	0.033 (2)	0.061 (3)	0.035 (2)	-0.001 (2)	-0.0091 (19)	-0.003 (2)
N2	0.045 (3)	0.058 (3)	0.035 (2)	0.000 (2)	-0.009 (2)	-0.002 (2)
C1	0.035 (2)	0.049 (3)	0.033 (2)	-0.004 (2)	0.001 (2)	0.003 (2)
C2	0.046 (3)	0.058 (3)	0.055 (4)	0.011 (2)	0.003 (3)	0.006 (3)
C3	0.067 (4)	0.059 (3)	0.063 (4)	0.002 (3)	0.008 (3)	-0.008 (3)
C4	0.061 (3)	0.062 (3)	0.041 (3)	-0.013 (3)	0.005 (3)	-0.007 (3)
C5	0.052 (3)	0.066 (3)	0.040 (3)	-0.001 (3)	-0.013 (2)	-0.001 (3)
C6	0.038 (2)	0.055 (3)	0.041 (2)	0.0026 (19)	-0.005 (3)	-0.003 (3)
C7	0.097 (4)	0.095 (5)	0.058 (3)	-0.016 (6)	-0.009 (3)	-0.021 (5)
C8	0.029 (2)	0.059 (3)	0.040 (3)	-0.003 (2)	0.000 (2)	0.008 (2)
C9	0.049 (3)	0.046 (3)	0.039 (3)	-0.003 (2)	-0.013 (2)	-0.002 (2)
C10	0.057 (3)	0.041 (2)	0.033 (2)	-0.008 (2)	-0.006 (3)	-0.002 (3)
C11	0.070 (4)	0.045 (3)	0.043 (3)	-0.014 (3)	0.000 (3)	-0.002 (2)
C12	0.116 (6)	0.059 (4)	0.042 (4)	-0.021 (4)	0.016 (4)	-0.006 (3)
C13	0.152 (7)	0.059 (4)	0.034 (3)	-0.030 (4)	-0.021 (4)	-0.003 (3)
C14	0.117 (6)	0.082 (5)	0.050 (4)	-0.001 (5)	-0.043 (4)	0.002 (4)
C15	0.075 (4)	0.054 (3)	0.053 (3)	-0.002 (3)	-0.020 (3)	0.003 (3)

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

Br1—C11	1.891 (6)	C6—H6	0.9300
O1—C8	1.225 (6)	C7—H7C	0.9600
N1—C8	1.351 (6)	C7—H7A	0.9600
N1—N2	1.379 (5)	C7—H7B	0.9600
N1—H1	0.837 (19)	C9—C10	1.470 (8)
N2—C9	1.270 (6)	C9—H9	0.9300
C1—C2	1.375 (7)	C10—C11	1.381 (7)
C1—C6	1.389 (6)	C10—C15	1.386 (7)
C1—C8	1.485 (6)	C11—C12	1.385 (7)
C2—C3	1.375 (8)	C12—C13	1.378 (10)
C2—H2	0.9300	C12—H12	0.9300
C3—C4	1.371 (8)	C13—C14	1.340 (10)
C3—H3	0.9300	C13—H13	0.9300
C4—C5	1.386 (7)	C14—C15	1.370 (8)
C4—C7	1.508 (9)	C14—H14	0.9300
C5—C6	1.384 (8)	C15—H15	0.9300
C5—H5	0.9300		
C8—N1—N2	119.2 (4)	H7C—C7—H7B	109.5
C8—N1—H1	123 (4)	H7A—C7—H7B	109.5
N2—N1—H1	117 (4)	O1—C8—N1	121.7 (4)
C9—N2—N1	116.0 (4)	O1—C8—C1	121.6 (4)
C2—C1—C6	118.6 (5)	N1—C8—C1	116.6 (4)

C2—C1—C8	117.6 (4)	N2—C9—C10	120.1 (4)
C6—C1—C8	123.8 (5)	N2—C9—H9	120.0
C1—C2—C3	121.0 (5)	C10—C9—H9	120.0
C1—C2—H2	119.5	C11—C10—C15	118.1 (5)
C3—C2—H2	119.5	C11—C10—C9	122.5 (5)
C4—C3—C2	121.2 (5)	C15—C10—C9	119.4 (5)
C4—C3—H3	119.4	C10—C11—C12	120.5 (6)
C2—C3—H3	119.4	C10—C11—Br1	122.2 (4)
C3—C4—C5	118.2 (5)	C12—C11—Br1	117.2 (5)
C3—C4—C7	121.8 (6)	C13—C12—C11	119.4 (6)
C5—C4—C7	120.0 (6)	C13—C12—H12	120.3
C6—C5—C4	121.1 (5)	C11—C12—H12	120.3
C6—C5—H5	119.5	C14—C13—C12	120.5 (6)
C4—C5—H5	119.5	C14—C13—H13	119.8
C5—C6—C1	120.0 (5)	C12—C13—H13	119.8
C5—C6—H6	120.0	C13—C14—C15	120.6 (7)
C1—C6—H6	120.0	C13—C14—H14	119.7
C4—C7—H7C	109.5	C15—C14—H14	119.7
C4—C7—H7A	109.5	C14—C15—C10	120.8 (6)
H7C—C7—H7A	109.5	C14—C15—H15	119.6
C4—C7—H7B	109.5	C10—C15—H15	119.6
C8—N1—N2—C9	-173.3 (5)	C6—C1—C8—N1	26.3 (7)
C6—C1—C2—C3	-1.5 (7)	N1—N2—C9—C10	175.7 (4)
C8—C1—C2—C3	178.4 (5)	N2—C9—C10—C11	161.4 (5)
C1—C2—C3—C4	1.0 (8)	N2—C9—C10—C15	-19.2 (7)
C2—C3—C4—C5	-0.2 (8)	C15—C10—C11—C12	-1.2 (7)
C2—C3—C4—C7	-179.2 (6)	C9—C10—C11—C12	178.2 (5)
C3—C4—C5—C6	-0.1 (8)	C15—C10—C11—Br1	178.1 (4)
C7—C4—C5—C6	178.9 (6)	C9—C10—C11—Br1	-2.5 (6)
C4—C5—C6—C1	-0.3 (8)	C10—C11—C12—C13	0.4 (8)
C2—C1—C6—C5	1.1 (7)	Br1—C11—C12—C13	-178.9 (5)
C8—C1—C6—C5	-178.8 (5)	C11—C12—C13—C14	-1.1 (10)
N2—N1—C8—O1	-3.5 (7)	C12—C13—C14—C15	2.8 (11)
N2—N1—C8—C1	174.6 (4)	C13—C14—C15—C10	-3.6 (10)
C2—C1—C8—O1	24.6 (7)	C11—C10—C15—C14	2.8 (8)
C6—C1—C8—O1	-155.5 (5)	C9—C10—C15—C14	-176.6 (6)
C2—C1—C8—N1	-153.6 (5)		

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N1—H1 \cdots O1 ⁱ	0.84 (3)	1.96 (3)	2.785 (5)	167 (5)
C9—H9 \cdots O1 ⁱ	0.93	2.38	3.166 (6)	142

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