N'-[(E)-2-Hydroxy-3,5-diiodobenzyl-idene]pyridine-3-carbohydrazide

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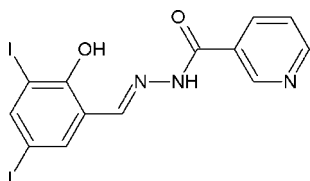Correspondence e-mail: rajagopal18@yahoo.com

In the title compound, $C_{13}H_9I_2N_3O_2$, the dihedral angle between the two aromatic rings is 10.5 (2)°. The molecule displays a *trans* configuration with respect to the C=N bond. An intramolecular O—H···N hydrogen bond occurs. The crystal packing is stabilized by N—H···O and C—H···O hydrogen bonds.

Related literature

For the biological activity of isoniazid derivatives, see: Janin (2007); Kahwa *et al.* (1986); Chen *et al.* (1997); Ren *et al.* (2002). For a related structure, see: Zhi & Wang (2010). For hydrogen-bond motifs, see: Bernstein *et al.* (1995).

Experimental

Crystal data

$C_{13}H_9I_2N_3O_2$	$a = 17.4800$ (5) Å
$M_r = 493.03$	$b = 8.5710$ (4) Å
Monoclinic, $P2_1/c$	$c = 9.8650$ (3) Å
$\beta = 90.451$ (4)°	$\mu = 4.26$ mm^{-1}
$V = 1477.94$ (9) Å^3	$T = 293$ K
$Z = 4$	$0.25 \times 0.22 \times 0.19$ mm
Mo $K\alpha$ radiation	

Data collection

Bruker APEXII CCD area-detector diffractometer	19735 measured reflections
Absorption correction: multi-scan (*SADABS*; Sheldrick, 2003) $T_{min} = 0.359$, $T_{max} = 0.445$	4866 independent reflections 3640 reflections with $I > 2\sigma(I)$ $R_{int} = 0.024$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.034$	183 parameters
$wR(F^2) = 0.106$	H-atom parameters constrained
$S = 0.99$	$\Delta\rho_{max} = 1.68$ e Å^{-3}
4866 reflections	$\Delta\rho_{min} = -1.17$ e Å^{-3}

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

D—H···A	D—H	H···A	D···A	D—H···A
O1—H1A···N1	0.82	1.88	2.599 (4)	146
N2—H2···O2^i	0.86	2.13	2.962 (4)	163
C7—H7···O2^i	0.93	2.59	3.367 (4)	142

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

Data collection: *APEX2* (Bruker, 2004); cell refinement: *SAINT* (Bruker, 2004); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3* (Farrugia, 1997); software used to prepare material for publication: *SHELXL97* and *PLATON* (Spek, 2009).

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supplementary materials

N'-[(*E*)-2-Hydroxy-3,5-diiodobenzylidene]pyridine-3-carbohydrazide

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Comment

In the search of new compounds, isoniazid derivatives have been found to possess potential tuberculostatic activity (Janin, 2007). Schiff bases have attracted much attention because of their biological activity (Kahwa *et al.*, 1986). Some of the compounds have been found to have excellent pharmacological and antibacterial activity (Chen *et al.*,1997; Ren *et al.*, 2002). Against this background, and in order to obtain detailed information on molecular conformations in the solid state, an X-ray study of the title compound was carried out.

X-Ray analysis confirms the molecular structure and atom connectivity as illustrated in Fig. 1. The pyridine ring is essentially planar with maximum deviations of 0.012 (5) Å, for atom C12. The molecular structure of the title compound displays a *trans* configuration with respect to the C=N and C—N bonds. The dihedral angle between the benzene and the pyridine rings is 10.5 (2)°. The atoms I1, I2 and O1 are deviated by -0.008 (1), -0.054 (1) and -0.050 (3) Å from the leastsquares plane of the benzene ring. All the bond lengths are within normal ranges and comparable to those in other similar compound (Zhi & Wang, 2010).

The intramolecular O1–H1A⋯N1 hydrogen bond completes a six-membered ring, which generates an S(6) motif (Bernstein *et al.*, 1995). Atoms N2 and C7 act as donors to form bifurcated hydrogen bonds with atom O2 as an acceptor, results in the formation of $R^2_1(6)$ bifurcated ring. In addition to van der Waals interaction, the crystal packing is stabilized N–H⋯O, O–H⋯N and C–H⋯O hydrogen bonds and further stabilized by weak intermolecular C2–I1⋯*Cg*2[ii] interaction involving the centroid of the benzene ring (*Cg*2 is the centroid of the benzene ring; ii= *x*,1/2 - *y*,-1/2 + *z*).

Experimental

An methanol solution (20 ml) and 3,5-diiodosalicylaldehyde (10 mmol) was magnetically stirred in a round bottom flask after getting clear solution followed by addition of Nicotinic hydrazide (10 mmol).The reaction mixture was refluxed for two hours and upon cooled to room temperature yellow crystalline solid was precipitated from the mixture it was recrystaliced with Dimethylformamide single yellow crystal were obtained washed with methanol and air dried. Single crystals suitable for X-ray diffraction were obtained by slow evaporation of a solution of the title compound in acetone at room temperature.

Refinement

All H atoms were fixed geometrically and allowed to ride on their parent atoms, with C—H distances fixed in the range 0.93–0.97 Å, N—H = 0.86Å and O—H = 0.82Å. U_{iso}(H) were set to 1.5U_{eq}(C) for methyl H and 1.2U_{eq}(C, N, O) for other H atoms.
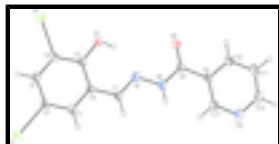
supplementary materials

Figures

Fig. 1. The structure of showing the atom-numbering scheme. The displacement ellipsoids are drawn at the 30% probability level. The disordered atoms are omitted for clarity.
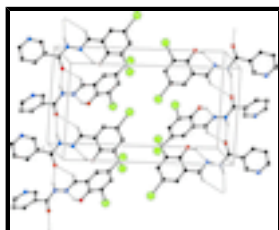

Fig. 2. A diagram showing the formation of $R^2_1(6)$ bifurcated ring and S(6) motif of molecules of the title compound through N–H···O, C–H···O and O–H···N hydrogen bonding.

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Crystal data

$C_{13}H_9I_2N_3O_2$	$F(000) = 920$
$M_r = 493.03$	$D_x = 2.216$ Mg m^{-3}
Monoclinic, $P2_1/c$	Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
Hall symbol: -P 2ybc	Cell parameters from 4867 reflections
$a = 17.4800$ (5) Å	$\theta = 1.2–31.6°$
$b = 8.5710$ (4) Å	$\mu = 4.26$ mm^{-1}
$c = 9.8650$ (3) Å	$T = 293$ K
$\beta = 90.451$ (4)°	Block, white
$V = 1477.94$ (9) Å^3	$0.25 \times 0.22 \times 0.19$ mm
$Z = 4$	

Data collection

Bruker APEXII CCD area-detector diffractometer	4866 independent reflections
Radiation source: fine-focus sealed tube	3640 reflections with $I > 2\sigma(I)$
graphite	$R_{int} = 0.024$
ω and φ scans	$\theta_{max} = 31.6°, \theta_{min} = 2.3°$
Absorption correction: multi-scan (*SADABS*; Sheldrick, 2003)	$h = -25 \rightarrow 25$
$T_{min} = 0.359, T_{max} = 0.445$	$k = -12 \rightarrow 12$
19735 measured reflections	$l = -14 \rightarrow 12$

Refinement

Refinement on F^2	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.034$	H-atom parameters constrained

$wR(F^2) = 0.106$	$w = 1/[\sigma^2(F_o^2) + (0.0525P)^2 + 1.9568P]$
	where $P = (F_o^2 + 2F_c^2)/3$
$S = 0.99$	$(\Delta/\sigma)_{max} = 0.001$
4866 reflections	$\Delta\rho_{max} = 1.68$ e Å^{-3}
183 parameters	$\Delta\rho_{min} = -1.17$ e Å^{-3}
0 restraints	Extinction correction: *SHELXL97* (Sheldrick, 2008), $Fc^* = kFc[1 + 0.001 x Fc^2 \lambda^3/\sin(2\theta)]^{-1/4}$
Primary atom site location: structure-invariant direct methods	Extinction coefficient: 0.0015 (2)

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 $F^2 > \sigma(F^2)$ 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 (Å^2)

	x	*y*	*z*	$U_{iso}*/U_{eq}$
C1	0.28906 (18)	0.3439 (4)	0.7602 (3)	0.0383 (7)
H1	0.2713	0.2819	0.6892	0.046*
C2	0.36332 (17)	0.3990 (4)	0.7591 (3)	0.0364 (6)
C3	0.39014 (17)	0.4933 (4)	0.8622 (3)	0.0367 (6)
H3	0.4397	0.5328	0.8598	0.044*
C4	0.34258 (17)	0.5285 (4)	0.9691 (3)	0.0371 (6)
C5	0.26788 (16)	0.4720 (4)	0.9750 (3)	0.0347 (6)
C6	0.24076 (16)	0.3806 (4)	0.8670 (3)	0.0343 (6)
C7	0.16298 (17)	0.3211 (4)	0.8627 (3)	0.0381 (7)
H7	0.1459	0.2645	0.7880	0.046*
C8	0.00141 (15)	0.2795 (4)	1.0619 (3)	0.0347 (6)
C9	−0.07420 (15)	0.2011 (4)	1.0435 (3)	0.0338 (6)
C10	−0.13285 (18)	0.2377 (5)	1.1314 (4)	0.0491 (9)
H10	−0.1259	0.3125	1.1987	0.059*
C11	−0.2021 (2)	0.1607 (5)	1.1169 (4)	0.0559 (10)
H11	−0.2425	0.1819	1.1748	0.067*
C12	−0.2097 (2)	0.0528 (5)	1.0153 (4)	0.0548 (9)
H12	−0.2569	0.0043	1.0046	0.066*
C13	−0.08758 (17)	0.0874 (4)	0.9470 (3)	0.0419 (7)
H13	−0.0477	0.0612	0.8894	0.050*
N1	0.11794 (15)	0.3469 (3)	0.9622 (3)	0.0394 (6)
N2	0.04524 (14)	0.2851 (4)	0.9506 (3)	0.0385 (6)
H2	0.0284	0.2511	0.8740	0.046*

N3	−0.15404 (18)	0.0128 (4)	0.9311 (4)	0.0546 (8)
O1	0.22488 (13)	0.5069 (3)	1.0828 (2)	0.0458 (6)
H1A	0.1816	0.4722	1.0714	0.069*
O2	0.02167 (13)	0.3328 (3)	1.1712 (2)	0.0481 (6)
I1	0.434960 (14)	0.34479 (3)	0.59728 (3)	0.05286 (10)
I2	0.385470 (15)	0.66245 (4)	1.12850 (3)	0.06771 (12)

Atomic displacement parameters (Å^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0354 (14)	0.0453 (18)	0.0344 (15)	−0.0064 (12)	0.0072 (11)	−0.0015 (12)
C2	0.0329 (13)	0.0427 (17)	0.0336 (14)	−0.0020 (12)	0.0085 (11)	0.0027 (12)
C3	0.0273 (12)	0.0402 (17)	0.0425 (16)	−0.0042 (11)	0.0051 (11)	0.0005 (13)
C4	0.0312 (13)	0.0416 (17)	0.0384 (15)	−0.0028 (11)	0.0036 (11)	−0.0044 (12)
C5	0.0301 (12)	0.0424 (17)	0.0318 (14)	−0.0012 (11)	0.0050 (10)	0.0014 (12)
C6	0.0268 (12)	0.0439 (17)	0.0324 (14)	−0.0055 (11)	0.0042 (10)	0.0026 (12)
C7	0.0304 (13)	0.0504 (19)	0.0336 (14)	−0.0077 (12)	0.0035 (11)	−0.0011 (13)
C8	0.0278 (12)	0.0457 (17)	0.0306 (13)	−0.0019 (11)	0.0026 (10)	0.0022 (12)
C9	0.0260 (12)	0.0463 (16)	0.0291 (13)	−0.0010 (11)	0.0023 (10)	0.0072 (12)
C10	0.0318 (14)	0.075 (3)	0.0405 (17)	−0.0020 (15)	0.0078 (12)	−0.0049 (17)
C11	0.0305 (15)	0.083 (3)	0.054 (2)	−0.0024 (16)	0.0129 (14)	0.0105 (19)
C12	0.0334 (16)	0.065 (2)	0.066 (2)	−0.0117 (15)	0.0036 (15)	0.010 (2)
C13	0.0314 (14)	0.053 (2)	0.0412 (17)	−0.0026 (13)	0.0049 (12)	0.0016 (14)
N1	0.0289 (11)	0.0539 (17)	0.0354 (13)	−0.0087 (10)	0.0033 (10)	0.0044 (11)
N2	0.0276 (11)	0.0598 (17)	0.0280 (11)	−0.0081 (11)	0.0039 (9)	0.0021 (11)
N3	0.0393 (15)	0.060 (2)	0.065 (2)	−0.0134 (13)	0.0004 (14)	−0.0036 (16)
O1	0.0350 (11)	0.0663 (17)	0.0363 (12)	−0.0061 (10)	0.0104 (9)	−0.0084 (11)
O2	0.0360 (11)	0.0756 (19)	0.0326 (12)	−0.0084 (11)	0.0010 (9)	−0.0064 (11)
I1	0.04408 (14)	0.06674 (19)	0.04806 (15)	−0.00641 (10)	0.01933 (10)	−0.00998 (11)
I2	0.04370 (14)	0.0942 (3)	0.06543 (19)	−0.02088 (13)	0.00969 (12)	−0.03975 (15)

Geometric parameters (Å, °)

C1—C2	1.382 (4)	C8—N2	1.344 (4)
C1—C6	1.391 (4)	C8—C9	1.492 (4)
C1—H1	0.9300	C9—C13	1.381 (5)
C2—C3	1.378 (4)	C9—C10	1.384 (4)
C2—I1	2.089 (3)	C10—C11	1.385 (5)
C3—C4	1.381 (4)	C10—H10	0.9300
C3—H3	0.9300	C11—C12	1.369 (6)
C4—C5	1.394 (4)	C11—H11	0.9300
C4—I2	2.082 (3)	C12—N3	1.329 (5)
C5—O1	1.341 (4)	C12—H12	0.9300
C5—C6	1.402 (4)	C13—N3	1.334 (4)
C6—C7	1.452 (4)	C13—H13	0.9300
C7—N1	1.282 (4)	N1—N2	1.381 (3)
C7—H7	0.9300	N2—H2	0.8600
C8—O2	1.221 (4)	O1—H1A	0.8200

C2—C1—C6	120.3 (3)	N2—C8—C9	115.3 (3)
C2—C1—H1	119.9	C13—C9—C10	118.0 (3)
C6—C1—H1	119.9	C13—C9—C8	123.1 (3)
C1—C2—C3	120.6 (3)	C10—C9—C8	118.8 (3)
C1—C2—I1	119.9 (2)	C11—C10—C9	118.6 (4)
C3—C2—I1	119.5 (2)	C11—C10—H10	120.7
C4—C3—C2	119.2 (3)	C9—C10—H10	120.7
C4—C3—H3	120.4	C12—C11—C10	118.5 (3)
C2—C3—H3	120.4	C12—C11—H11	120.7
C3—C4—C5	121.7 (3)	C10—C11—H11	120.7
C3—C4—I2	118.8 (2)	N3—C12—C11	124.3 (3)
C5—C4—I2	119.5 (2)	N3—C12—H12	117.8
O1—C5—C4	119.1 (3)	C11—C12—H12	117.8
O1—C5—C6	122.6 (3)	N3—C13—C9	124.2 (3)
C4—C5—C6	118.3 (3)	N3—C13—H13	117.9
C1—C6—C5	119.9 (3)	C9—C13—H13	117.9
C1—C6—C7	118.2 (3)	C7—N1—N2	116.2 (3)
C5—C6—C7	121.9 (3)	C8—N2—N1	118.5 (3)
N1—C7—C6	119.8 (3)	C8—N2—H2	120.8
N1—C7—H7	120.1	N1—N2—H2	120.8
C6—C7—H7	120.1	C12—N3—C13	116.4 (3)
O2—C8—N2	123.0 (3)	C5—O1—H1A	109.5
O2—C8—C9	121.8 (3)		
C6—C1—C2—C3	−1.2 (5)	C5—C6—C7—N1	−2.7 (5)
C6—C1—C2—I1	−179.4 (2)	O2—C8—C9—C13	−152.5 (3)
C1—C2—C3—C4	1.9 (5)	N2—C8—C9—C13	26.8 (5)
I1—C2—C3—C4	−179.8 (2)	O2—C8—C9—C10	24.1 (5)
C2—C3—C4—C5	−0.5 (5)	N2—C8—C9—C10	−156.6 (3)
C2—C3—C4—I2	177.2 (2)	C13—C9—C10—C11	−1.0 (5)
C3—C4—C5—O1	178.5 (3)	C8—C9—C10—C11	−177.8 (3)
I2—C4—C5—O1	0.8 (4)	C9—C10—C11—C12	−0.6 (6)
C3—C4—C5—C6	−1.6 (5)	C10—C11—C12—N3	2.1 (7)
I2—C4—C5—C6	−179.3 (2)	C10—C9—C13—N3	1.5 (5)
C2—C1—C6—C5	−1.0 (5)	C8—C9—C13—N3	178.1 (3)
C2—C1—C6—C7	179.4 (3)	C6—C7—N1—N2	−179.4 (3)
O1—C5—C6—C1	−177.7 (3)	O2—C8—N2—N1	3.4 (5)
C4—C5—C6—C1	2.3 (5)	C9—C8—N2—N1	−175.9 (3)
O1—C5—C6—C7	1.9 (5)	C7—N1—N2—C8	166.1 (3)
C4—C5—C6—C7	−178.0 (3)	C11—C12—N3—C13	−1.7 (6)
C1—C6—C7—N1	177.0 (3)	C9—C13—N3—C12	−0.2 (6)

Hydrogen-bond geometry (Å, °)

D—H⋯*A*	*D*—H	H⋯*A*	*D*⋯*A*	*D*—H⋯*A*
O1—H1A⋯N1	0.82	1.88	2.599 (4)	146
N2—H2⋯O2ⁱ	0.86	2.13	2.962 (4)	163.
C7—H7⋯O2ⁱ	0.93	2.59	3.367 (4)	142.

Symmetry codes: (i) *x*, −*y*+1/2, *z*−1/2.

Fig. 1

Fig. 2