

(E)-N'-(5-Chloro-2-hydroxybenzylidene)-2-nitrobenzohydrazide

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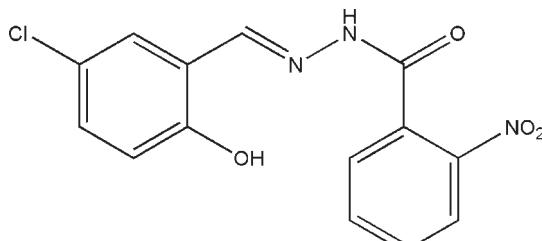
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Key indicators: single-crystal X-ray study; $T = 298\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$; R factor = 0.046; wR factor = 0.122; data-to-parameter ratio = 14.8.

In the title Schiff base compound, $\text{C}_{14}\text{H}_{10}\text{ClN}_3\text{O}_4$, the molecule adopts an *E* geometry with respect to the $\text{C}=\text{N}$ bond and an intramolecular $\text{O}-\text{H}\cdots\text{N}$ hydrogen bond is present. The benzene rings form a dihedral angle of $73.4(2)^\circ$. In the crystal, inversion dimers linked by pairs of $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds occur.

Related literature

For a related structure and background, see: Qian & Qu (2009).



Experimental

Crystal data

$\text{C}_{14}\text{H}_{10}\text{ClN}_3\text{O}_4$

$M_r = 319.70$

Triclinic, $P\bar{1}$	$V = 704.9(2)\text{ \AA}^3$
$a = 7.353(1)\text{ \AA}$	$Z = 2$
$b = 10.005(2)\text{ \AA}$	Mo $K\alpha$ radiation
$c = 10.273(2)\text{ \AA}$	$\mu = 0.29\text{ mm}^{-1}$
$\alpha = 93.393(3)^\circ$	$T = 298\text{ K}$
$\beta = 108.144(3)^\circ$	$0.23 \times 0.20 \times 0.20\text{ mm}$
$\gamma = 98.886(4)^\circ$	

Data collection

Bruker SMART CCD	4324 measured reflections
diffractometer	3004 independent reflections
Absorption correction: multi-scan (<i>SADABS</i> ; Bruker, 2001)	2237 reflections with $I > 2\sigma(I)$
$T_{\min} = 0.936$, $T_{\max} = 0.944$	$R_{\text{int}} = 0.013$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.046$	H atoms treated by a mixture of independent and constrained refinement
$wR(F^2) = 0.122$	$\Delta\rho_{\max} = 0.30\text{ e \AA}^{-3}$
$S = 1.02$	$\Delta\rho_{\min} = -0.44\text{ e \AA}^{-3}$
3004 reflections	
203 parameters	
1 restraint	

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

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
O1—H1 \cdots N1	0.82	1.94	2.657 (2)	145
N2—H2 \cdots O2 ⁱ	0.902 (10)	1.966 (11)	2.863 (2)	173 (3)

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

Data collection: *SMART* (Bruker, 2007); cell refinement: *SAINT* (Bruker, 2007); data reduction: *SAINT*; program(s) used to solve structure: *SHELXTL* (Sheldrick, 2008); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB5055).

References

- Bruker (2001). *SADABS*. Bruker AXS Inc., Madison, Wisconsin, USA.
Bruker (2007). *SMART* and *SAINT*. Bruker AXS Inc., Madison, Wisconsin, USA.
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Sheldrick, G. M. (2008). *Acta Cryst. A64*, 112–122.

supporting information

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(E)-N'-(5-Chloro-2-hydroxybenzylidene)-2-nitrobenzohydrazide

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S1. Comment

As part of our ongoing studies of Schiff bases (Qian & Qu, 2009), we now report the synthesis and structure of the title compound, (I), (Fig. 1).

In the title compound, the Schiff base molecule adopts an *E* geometry with respect to the CN bond, as shown in Fig. 1. There forms an intramolecular O—H···N hydrogen bond. The two benzene rings form a dihedral angle of 73.4 (2)°. The dihedral angle between the O3/N3/O4 plane and the C9—C14 benzene ring is 23.2 (2)°. In the crystal structure, the adjacent two Schiff base molecules are linked through intermolecular N—H···O hydrogen bonds (Table 1) to form a dimer (Fig. 2).

S2. Experimental

2-Nitrobenzohydrazide (1 mmol, 0.181 g) and 5-chlorosalicylaldehyde (1 mmol, 0.156 g) were dissolved in anhydrous methanol (15 ml). The mixture was stirred for several minutes at room temperature. The product was isolated and recrystallized from methanol, colorless blocks of (I) were obtained after a week.

S3. Refinement

The imino H atom was located in a difference map and its positional parameters were refined with a fixed isotropic thermal parameter of 0.08 Å². Other H atoms were positioned geometrically and refined as riding with C—H = 0.93 Å, O—H = 0.82 Å, and with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ and $1.5U_{\text{eq}}(\text{O})$.

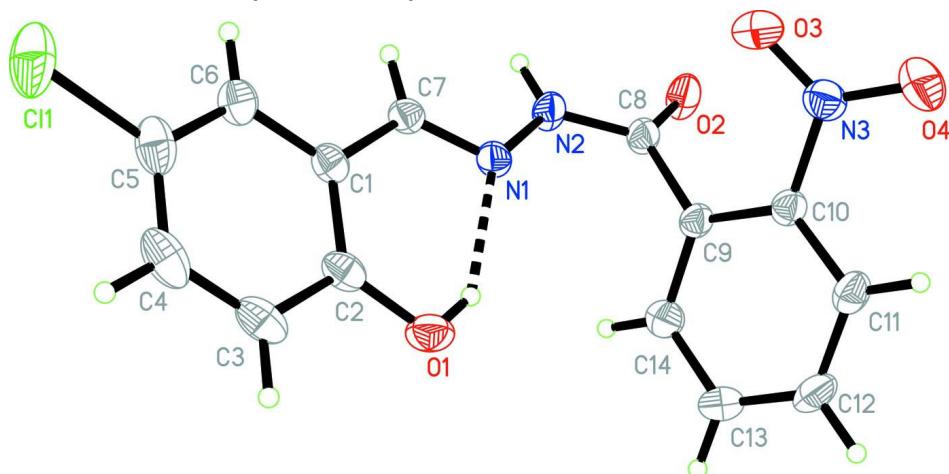
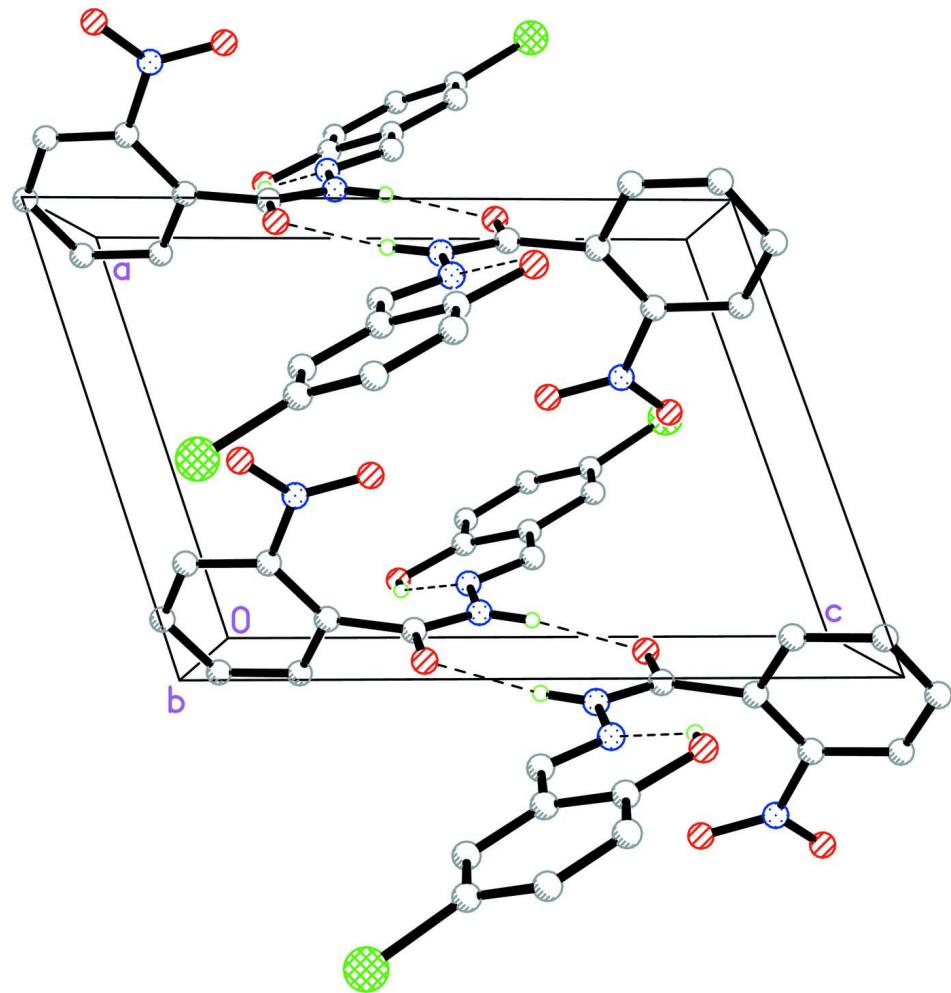


Figure 1

The molecular structure of (I). Displacement ellipsoids are drawn at the 30% probability level. Hydrogen bonding is shown by dashed lines.

**Figure 2**

The molecular packing of (I), viewed along the *b* axis. Hydrogen bonding is shown in dashed lines.

(*E*)-*N'*-(5-Chloro-2-hydroxybenzylidene)-2-nitrobenzohydrazide

Crystal data

$C_{14}H_{10}ClN_3O_4$

$M_r = 319.70$

Triclinic, $P\bar{1}$

Hall symbol: -P 1

$a = 7.353 (1) \text{ \AA}$

$b = 10.005 (2) \text{ \AA}$

$c = 10.273 (2) \text{ \AA}$

$\alpha = 93.393 (3)^\circ$

$\beta = 108.144 (3)^\circ$

$\gamma = 98.886 (4)^\circ$

$V = 704.9 (2) \text{ \AA}^3$

$Z = 2$

$F(000) = 328$

$D_x = 1.506 \text{ Mg m}^{-3}$

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

Cell parameters from 1356 reflections

$\theta = 2.4\text{--}25.6^\circ$

$\mu = 0.29 \text{ mm}^{-1}$

$T = 298 \text{ K}$

Block, colorless

$0.23 \times 0.20 \times 0.20 \text{ mm}$

Data collection

Bruker SMART CCD
diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
 ω scans
Absorption correction: multi-scan
(*SADABS*; Bruker, 2001)
 $T_{\min} = 0.936$, $T_{\max} = 0.944$

4324 measured reflections
3004 independent reflections
2237 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.013$
 $\theta_{\max} = 27.0^\circ$, $\theta_{\min} = 2.1^\circ$
 $h = -9 \rightarrow 9$
 $k = -9 \rightarrow 12$
 $l = -13 \rightarrow 12$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.046$
 $wR(F^2) = 0.122$
 $S = 1.02$
3004 reflections
203 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
 $w = 1/[\sigma^2(F_o^2) + (0.0501P)^2 + 0.3008P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.30 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.44 \text{ e } \text{\AA}^{-3}$

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 (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
C11	0.54671 (11)	-0.14057 (9)	0.86494 (8)	0.0837 (3)
N1	0.1470 (2)	0.23302 (15)	0.43169 (16)	0.0372 (4)
N2	0.0876 (3)	0.35676 (17)	0.43461 (17)	0.0422 (4)
N3	0.3643 (3)	0.53956 (19)	0.2180 (2)	0.0510 (5)
O1	0.1405 (2)	-0.01201 (16)	0.31361 (15)	0.0534 (4)
H1	0.1202	0.0663	0.3150	0.080*
O2	-0.0068 (3)	0.53476 (15)	0.32927 (15)	0.0574 (5)
O3	0.4103 (3)	0.54714 (19)	0.34314 (18)	0.0735 (5)
O4	0.4446 (3)	0.6178 (2)	0.1575 (2)	0.0898 (7)
C1	0.2638 (3)	0.05581 (19)	0.5597 (2)	0.0383 (4)
C2	0.2290 (3)	-0.0378 (2)	0.4431 (2)	0.0428 (5)
C3	0.2854 (3)	-0.1637 (2)	0.4603 (3)	0.0543 (6)
H3	0.2585	-0.2270	0.3837	0.065*
C4	0.3798 (4)	-0.1956 (2)	0.5886 (3)	0.0615 (7)
H4	0.4185	-0.2796	0.5990	0.074*

C5	0.4176 (3)	-0.1023 (3)	0.7029 (3)	0.0551 (6)
C6	0.3583 (3)	0.0212 (2)	0.6893 (2)	0.0463 (5)
H6	0.3816	0.0820	0.7673	0.056*
C7	0.2062 (3)	0.18811 (19)	0.5484 (2)	0.0377 (4)
H7	0.2125	0.2412	0.6277	0.045*
C8	0.0525 (3)	0.4269 (2)	0.3254 (2)	0.0404 (5)
C9	0.0740 (3)	0.36796 (19)	0.19420 (19)	0.0369 (4)
C10	0.2071 (3)	0.43017 (19)	0.13654 (19)	0.0374 (4)
C11	0.2035 (3)	0.3885 (2)	0.0051 (2)	0.0461 (5)
H11	0.2919	0.4344	-0.0323	0.055*
C12	0.0674 (4)	0.2784 (2)	-0.0695 (2)	0.0529 (6)
H12	0.0638	0.2488	-0.1580	0.063*
C13	-0.0624 (4)	0.2123 (2)	-0.0142 (2)	0.0573 (6)
H13	-0.1526	0.1366	-0.0643	0.069*
C14	-0.0606 (4)	0.2574 (2)	0.1164 (2)	0.0516 (6)
H14	-0.1516	0.2126	0.1522	0.062*
H2	0.072 (4)	0.392 (3)	0.5126 (18)	0.080*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C11	0.0709 (5)	0.0970 (6)	0.0952 (6)	0.0360 (4)	0.0260 (4)	0.0547 (5)
N1	0.0494 (10)	0.0296 (8)	0.0381 (9)	0.0096 (7)	0.0212 (8)	0.0020 (7)
N2	0.0654 (11)	0.0356 (9)	0.0368 (9)	0.0178 (8)	0.0281 (8)	0.0055 (7)
N3	0.0529 (11)	0.0467 (11)	0.0539 (12)	0.0027 (9)	0.0224 (9)	-0.0015 (9)
O1	0.0639 (10)	0.0449 (9)	0.0481 (9)	0.0110 (8)	0.0156 (8)	-0.0083 (7)
O2	0.1017 (13)	0.0428 (9)	0.0476 (9)	0.0330 (9)	0.0414 (9)	0.0136 (7)
O3	0.0818 (13)	0.0720 (12)	0.0495 (10)	-0.0114 (10)	0.0125 (9)	-0.0109 (9)
O4	0.0946 (15)	0.0820 (14)	0.0882 (14)	-0.0283 (12)	0.0452 (12)	0.0039 (11)
C1	0.0381 (10)	0.0339 (10)	0.0461 (11)	0.0053 (8)	0.0189 (9)	0.0053 (8)
C2	0.0377 (11)	0.0351 (10)	0.0571 (13)	0.0023 (8)	0.0205 (10)	-0.0005 (9)
C3	0.0524 (13)	0.0341 (11)	0.0804 (17)	0.0077 (10)	0.0291 (13)	-0.0022 (11)
C4	0.0554 (14)	0.0390 (12)	0.103 (2)	0.0170 (11)	0.0389 (15)	0.0186 (13)
C5	0.0443 (12)	0.0553 (14)	0.0752 (16)	0.0167 (10)	0.0253 (12)	0.0291 (12)
C6	0.0459 (12)	0.0451 (12)	0.0527 (12)	0.0102 (9)	0.0207 (10)	0.0112 (10)
C7	0.0441 (11)	0.0336 (10)	0.0386 (10)	0.0053 (8)	0.0194 (9)	0.0001 (8)
C8	0.0553 (12)	0.0350 (10)	0.0369 (10)	0.0098 (9)	0.0227 (9)	0.0046 (8)
C9	0.0488 (11)	0.0337 (10)	0.0322 (9)	0.0115 (8)	0.0168 (8)	0.0035 (8)
C10	0.0444 (11)	0.0355 (10)	0.0349 (10)	0.0108 (8)	0.0150 (9)	0.0034 (8)
C11	0.0558 (13)	0.0521 (13)	0.0388 (11)	0.0154 (10)	0.0242 (10)	0.0080 (9)
C12	0.0706 (16)	0.0597 (14)	0.0300 (10)	0.0190 (12)	0.0163 (10)	-0.0004 (10)
C13	0.0663 (15)	0.0525 (14)	0.0424 (12)	-0.0003 (11)	0.0103 (11)	-0.0083 (10)
C14	0.0580 (14)	0.0488 (13)	0.0476 (12)	-0.0015 (10)	0.0229 (11)	-0.0004 (10)

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

C11—C5	1.738 (3)	C4—C5	1.383 (4)
N1—C7	1.276 (2)	C4—H4	0.9300

N1—N2	1.377 (2)	C5—C6	1.373 (3)
N2—C8	1.336 (2)	C6—H6	0.9300
N2—H2	0.902 (10)	C7—H7	0.9300
N3—O3	1.217 (2)	C8—C9	1.503 (2)
N3—O4	1.221 (2)	C9—C14	1.380 (3)
N3—C10	1.460 (3)	C9—C10	1.383 (3)
O1—C2	1.347 (3)	C10—C11	1.380 (3)
O1—H1	0.8200	C11—C12	1.373 (3)
O2—C8	1.228 (2)	C11—H11	0.9300
C1—C6	1.389 (3)	C12—C13	1.364 (3)
C1—C2	1.407 (3)	C12—H12	0.9300
C1—C7	1.451 (3)	C13—C14	1.385 (3)
C2—C3	1.390 (3)	C13—H13	0.9300
C3—C4	1.367 (4)	C14—H14	0.9300
C3—H3	0.9300		
C7—N1—N2	115.34 (15)	C1—C6—H6	119.8
C8—N2—N1	122.02 (15)	N1—C7—C1	121.21 (17)
C8—N2—H2	118.8 (18)	N1—C7—H7	119.4
N1—N2—H2	119.2 (18)	C1—C7—H7	119.4
O3—N3—O4	123.3 (2)	O2—C8—N2	120.66 (17)
O3—N3—C10	118.26 (18)	O2—C8—C9	120.00 (17)
O4—N3—C10	118.47 (19)	N2—C8—C9	119.22 (17)
C2—O1—H1	109.5	C14—C9—C10	117.09 (18)
C6—C1—C2	118.97 (19)	C14—C9—C8	119.38 (18)
C6—C1—C7	119.03 (18)	C10—C9—C8	122.92 (18)
C2—C1—C7	121.99 (18)	C11—C10—C9	122.36 (19)
O1—C2—C3	117.59 (19)	C11—C10—N3	117.72 (18)
O1—C2—C1	122.93 (18)	C9—C10—N3	119.85 (17)
C3—C2—C1	119.5 (2)	C12—C11—C10	119.0 (2)
C4—C3—C2	120.7 (2)	C12—C11—H11	120.5
C4—C3—H3	119.6	C10—C11—H11	120.5
C2—C3—H3	119.6	C13—C12—C11	120.11 (19)
C3—C4—C5	119.7 (2)	C13—C12—H12	119.9
C3—C4—H4	120.1	C11—C12—H12	119.9
C5—C4—H4	120.1	C12—C13—C14	120.3 (2)
C6—C5—C4	120.7 (2)	C12—C13—H13	119.9
C6—C5—Cl1	119.7 (2)	C14—C13—H13	119.9
C4—C5—Cl1	119.51 (19)	C9—C14—C13	121.1 (2)
C5—C6—C1	120.3 (2)	C9—C14—H14	119.4
C5—C6—H6	119.8	C13—C14—H14	119.4

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H···A
O1—H1···N1	0.82	1.94	2.657 (2)	145

supporting information

N2—H2···O2 ⁱ	0.90 (1)	1.97 (1)	2.863 (2)	173 (3)
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Symmetry code: (i) $-x, -y+1, -z+1$.