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(*E*)-*N*′-(2-Chlorobenzylidene)-3,5-dihydroxybenzohydrazide dihydrate

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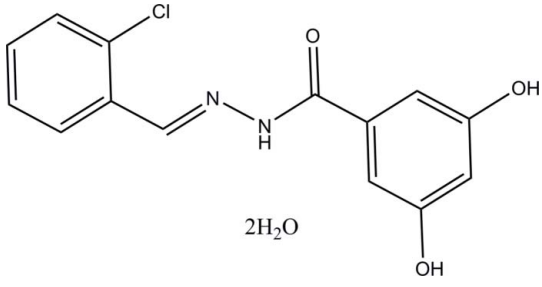
Key indicators: single-crystal X-ray study; T = 296 K; mean σ(C–C) = 0.004 Å; R factor = 0.051; wR factor = 0.140; data-to-parameter ratio = 17.7.

In the Schiff base molecule of the title compound, $C_{14}H_{11}ClN_2O_3\cdot 2H_2O$, the benzene rings form a dihedral angle of 20.6 (1)°. The water molecules of crystallization are involved in the formation of a three-dimensional hydrogen-bonding network *via* O—H···O and N—H···O hydrogen bonds.

Related literature

For general background to Schiff base compounds, see: Brückner *et al.* (2000); Harrop *et al.*(2003); Zhang *et al.* (2008). For related structures, see: Diao *et al.* (2007); Jiang *et al.* (2008); Huang *et al.* (2008); Deng *et al.* (2009).



Experimental

Crystal data

$C_{14}H_{11}ClN_2O_3\cdot 2H_2O$
M_r = 326.73
Monoclinic, $P2_1/c$
a = 8.023 (2) Å
b = 11.852 (4) Å
c = 16.318 (5) Å
β = 100.387 (4)°
V = 1526.1 (8) Å^3
Z = 4
Mo $K\alpha$ radiation
μ = 0.28 mm^{-1}
T = 296 K
0.44 × 0.12 × 0.07 mm

Data collection

Bruker APEXII CCD diffractometer
12729 measured reflections
3522 independent reflections
2286 reflections with $I > 2\sigma(I)$
R_{int} = 0.040

Refinement

$R[F^2 > 2\sigma(F^2)]$ = 0.051
$wR(F^2)$ = 0.140
S = 1.03
3522 reflections
199 parameters
H-atom parameters constrained
$\Delta\rho_{max}$ = 0.32 e Å^{-3}
$\Delta\rho_{min}$ = −0.33 e Å^{-3}

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

D—H···A	D—H	H···A	D···A	D—H···A
O2—H2*B*···O5	0.82	1.87	2.674 (3)	168
O3—H3*A*···O1^{i}	0.82	1.85	2.665 (2)	169
N2—H2*A*···O2^{ii}	0.86	2.36	3.196 (3)	164
O4—H4*B*···O1	0.85	2.12	2.960 (4)	171
O5—H5*A*···O4^{iii}	0.85	1.99	2.816 (4)	163
O5—H5*B*···O3^{iv}	0.85	2.14	2.902 (3)	150

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

Data collection: *APEX2* (Bruker, 2005); cell refinement: *APEX2*; data reduction: *SAINT* (Bruker, 2005); 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: *SHELXL97*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: CV5172).

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(*E*)-*N*′-(2-Chlorobenzylidene)-3,5-dihydroxybenzohydrazide dihydrate

Ling Yuan, Yi Nan, Jing-Yuan Li and Xiu-Lan Huang

S1. Comment

Schiff base compounds are known to exhibit antibacterial and antitumor properties (Brückner *et al.*, 2000; Harrop *et al.*, 2003; Zhang *et al.*, 2008). In order to expand this filed, we report here the structure of the title compound (I).

In (I) (Fig. 1), all bond lengths and angles are normal and comparable with those found in the related compounds (Diao *et al.*, 2007; Deng *et al.*, 2009; Huang *et al.*, 2008, Jiang *et al.*, 2008). In the Shiff base molecule, two benzene rings form a dihedral angle of 20.6 (1)°.

In the crystal structure, intermolecular O—H···O and N—H···O hydrogen bonds (Table 1) consolidate the crystal packing.

S2. Experimental

2-Chlorobenzaldehyde (0.1 mmol, 14.1 mg) and 3,5-dihydroxybenzhydrazide (0.1 mmol, 16.8 mg) were dissolved in a methanol solution (10 ml). The mixture was stirred at room temperature for 1 h and filtered. After keeping the filtrate in air for three days, yellow crystals were formed.

S3. Refinement

H atoms were placed in calculated positions (C—H 0.93 Å; N—H 0.86 Å; O—H 0.82 Å) and were included in the refinement in the riding model approximation, with U_{iso}(H)=) =1.2-1.5U_{eq} of the parent atom.

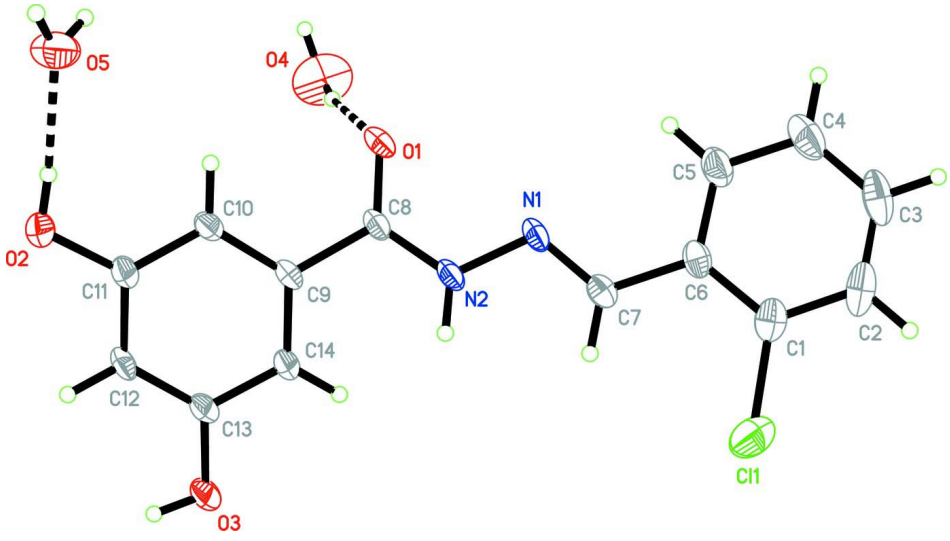

Figure 1

The molecular structure of (I) showing the atomic numbering and 30% probability displacement ellipsoids.

(*E*)-*N*′-(2-Chlorobenzylidene)-3,5-dihydroxybenzohydrazide dihydrate

Crystal data

$C_{14}H_{11}ClN_2O_3 \cdot 2H_2O$
$M_r = 326.73$
Monoclinic, $P2_1/c$
Hall symbol: -P2ybc
$a = 8.023$ (2) Å
$b = 11.852$ (4) Å
$c = 16.318$ (5) Å
$\beta = 100.387$ (4)°
$V = 1526.1$ (8) Å^3
$Z = 4$
$F(000) = 680$
$D_x = 1.422$ Mg m^{-3}
Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
Cell parameters from 4061 reflections
$\theta = 2.5$–$27.1°$
$\mu = 0.28$ mm^{-1}
$T = 296$ K
Stick, yellow
0.44 × 0.12 × 0.07 mm

Data collection

Bruker APEXII CCD diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
φ and ω scans
12729 measured reflections
3522 independent reflections
2286 reflections with $I > 2\sigma(I)$
$R_{int} = 0.040$
$\theta_{max} = 27.6°$, $\theta_{min} = 2.1°$
$h = -10 \rightarrow 10$
$k = -15 \rightarrow 15$
$l = -20 \rightarrow 21$

Refinement

Refinement on F^2
Least-squares matrix: full
$R[F^2 > 2\sigma(F^2)] = 0.051$
$wR(F^2) = 0.140$
$S = 1.03$
3522 reflections
199 parameters
0 restraints
Primary atom site location: structure-invariant direct methods
Secondary atom site location: difference Fourier map
Hydrogen site location: inferred from neighbouring sites
H-atom parameters constrained
$w = 1/[\sigma^2(F_o^2) + (0.044P)^2 + 1.2P]$ where $P = (F_o^2 + 2F_c^2)/3$
$(\Delta/\sigma)_{max} < 0.001$
$\Delta\rho_{max} = 0.32$ e Å^{-3}
$\Delta\rho_{min} = -0.33$ e Å^{-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 (Å^2)

	x	*y*	*z*	U_{iso}*/U_{eq}
Cl1	0.07549 (13)	0.73732 (7)	0.46696 (6)	0.0836 (3)
O1	0.4482 (3)	0.20583 (16)	0.45642 (10)	0.0555 (5)
O2	0.7564 (2)	0.07180 (15)	0.22642 (10)	0.0508 (5)
H2B	0.7932	0.0426	0.2716	0.076*
O3	0.3480 (3)	0.33745 (18)	0.10033 (10)	0.0615 (6)

H3A	0.3864	0.3177	0.0593	0.092*
N1	0.3161 (2)	0.41028 (17)	0.48357 (10)	0.0374 (4)
N2	0.3589 (2)	0.37857 (17)	0.40850 (10)	0.0368 (4)
H2A	0.3450	0.4252	0.3674	0.044*
C1	0.1081 (3)	0.6554 (2)	0.55602 (17)	0.0472 (6)
C2	0.0554 (4)	0.6968 (3)	0.6271 (2)	0.0621 (8)
H2	0.0022	0.7666	0.6257	0.074*
C3	0.0818 (4)	0.6351 (3)	0.6986 (2)	0.0678 (9)
H3	0.0483	0.6635	0.7462	0.081*
C4	0.1571 (4)	0.5318 (3)	0.70062 (18)	0.0687 (9)
H4	0.1738	0.4897	0.7495	0.082*
C5	0.2090 (4)	0.4891 (3)	0.63044 (16)	0.0550 (7)
H5	0.2596	0.4184	0.6326	0.066*
C6	0.1866 (3)	0.5505 (2)	0.55659 (14)	0.0382 (5)
C7	0.2396 (3)	0.5047 (2)	0.48164 (14)	0.0371 (5)
H7	0.2171	0.5453	0.4321	0.044*
C8	0.4220 (3)	0.2757 (2)	0.39941 (12)	0.0353 (5)
C9	0.4635 (3)	0.25014 (19)	0.31552 (12)	0.0326 (5)
C10	0.5859 (3)	0.16929 (19)	0.31167 (12)	0.0347 (5)
H10	0.6357	0.1302	0.3592	0.042*
C11	0.6334 (3)	0.14750 (19)	0.23525 (13)	0.0351 (5)
C12	0.5573 (3)	0.2046 (2)	0.16457 (12)	0.0387 (5)
H12	0.5916	0.1910	0.1140	0.046*
C13	0.4305 (3)	0.2818 (2)	0.16880 (12)	0.0387 (5)
C14	0.3828 (3)	0.3059 (2)	0.24457 (13)	0.0393 (5)
H14	0.2981	0.3585	0.2477	0.047*
O4	0.2697 (4)	−0.0135 (3)	0.4316 (2)	0.1159 (11)
H4A	0.3177	−0.0688	0.4595	0.174*
H4B	0.3266	0.0473	0.4339	0.174*
O5	0.9176 (3)	−0.00817 (18)	0.37266 (13)	0.0690 (6)
H5A	1.0182	−0.0070	0.4001	0.104*
H5B	0.8554	−0.0465	0.3993	0.104*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Cl1	0.1051 (7)	0.0521 (5)	0.0969 (7)	0.0157 (5)	0.0274 (6)	0.0134 (4)
O1	0.0881 (14)	0.0572 (11)	0.0272 (8)	0.0154 (10)	0.0261 (9)	0.0084 (8)
O2	0.0624 (12)	0.0562 (11)	0.0360 (9)	0.0213 (9)	0.0146 (8)	−0.0016 (8)
O3	0.0820 (14)	0.0817 (14)	0.0224 (8)	0.0345 (12)	0.0139 (8)	0.0076 (8)
N1	0.0404 (11)	0.0505 (12)	0.0244 (8)	0.0010 (9)	0.0141 (8)	−0.0046 (8)
N2	0.0438 (11)	0.0479 (11)	0.0217 (8)	0.0006 (9)	0.0140 (8)	−0.0011 (8)
C1	0.0379 (14)	0.0438 (14)	0.0619 (16)	−0.0085 (11)	0.0146 (12)	−0.0153 (12)
C2	0.0455 (16)	0.0534 (17)	0.093 (2)	−0.0089 (13)	0.0284 (16)	−0.0352 (17)
C3	0.0585 (19)	0.087 (2)	0.067 (2)	−0.0154 (17)	0.0349 (16)	−0.0362 (18)
C4	0.071 (2)	0.097 (3)	0.0449 (15)	0.0004 (19)	0.0279 (14)	−0.0074 (16)
C5	0.0568 (17)	0.0714 (19)	0.0414 (14)	0.0092 (14)	0.0212 (12)	−0.0046 (13)
C6	0.0311 (12)	0.0477 (13)	0.0384 (12)	−0.0053 (10)	0.0131 (9)	−0.0095 (10)

C7	0.0366 (12)	0.0455 (13)	0.0311 (11)	−0.0054 (11)	0.0114 (9)	−0.0022 (9)
C8	0.0394 (12)	0.0451 (13)	0.0240 (10)	0.0002 (10)	0.0129 (9)	−0.0016 (9)
C9	0.0368 (12)	0.0419 (12)	0.0209 (9)	−0.0035 (10)	0.0103 (8)	−0.0017 (8)
C10	0.0403 (13)	0.0412 (12)	0.0235 (9)	0.0004 (10)	0.0081 (9)	0.0006 (9)
C11	0.0406 (13)	0.0376 (12)	0.0290 (10)	−0.0009 (10)	0.0112 (9)	−0.0055 (9)
C12	0.0497 (14)	0.0474 (13)	0.0220 (9)	0.0015 (11)	0.0146 (9)	−0.0062 (9)
C13	0.0464 (13)	0.0481 (13)	0.0222 (10)	0.0042 (11)	0.0076 (9)	0.0001 (9)
C14	0.0464 (14)	0.0480 (13)	0.0257 (10)	0.0100 (11)	0.0124 (9)	−0.0005 (9)
O4	0.099 (2)	0.098 (2)	0.138 (3)	−0.0265 (18)	−0.0111 (19)	0.0009 (19)
O5	0.0685 (14)	0.0749 (14)	0.0616 (12)	0.0020 (11)	0.0063 (10)	0.0216 (11)

Geometric parameters (Å, º)

Cl1—C1	1.728 (3)	C5—C6	1.391 (4)
O1—C8	1.235 (3)	C5—H5	0.9300
O2—C11	1.361 (3)	C6—C7	1.469 (3)
O2—H2B	0.8200	C7—H7	0.9300
O3—C13	1.362 (3)	C8—C9	1.497 (3)
O3—H3A	0.8200	C9—C10	1.382 (3)
N1—C7	1.274 (3)	C9—C14	1.388 (3)
N1—N2	1.383 (2)	C10—C11	1.392 (3)
N2—C8	1.338 (3)	C10—H10	0.9300
N2—H2A	0.8600	C11—C12	1.381 (3)
C1—C6	1.392 (4)	C12—C13	1.379 (3)
C1—C2	1.393 (4)	C12—H12	0.9300
C2—C3	1.361 (5)	C13—C14	1.388 (3)
C2—H2	0.9300	C14—H14	0.9300
C3—C4	1.363 (5)	O4—H4A	0.8500
C3—H3	0.9300	O4—H4B	0.8500
C4—C5	1.383 (4)	O5—H5A	0.8500
C4—H4	0.9300	O5—H5B	0.8500
C11—O2—H2B	109.5	N1—C7—H7	119.5
C13—O3—H3A	109.5	C6—C7—H7	119.5
C7—N1—N2	114.36 (19)	O1—C8—N2	122.96 (18)
C8—N2—N1	120.34 (18)	O1—C8—C9	121.1 (2)
C8—N2—H2A	119.8	N2—C8—C9	115.90 (19)
N1—N2—H2A	119.8	C10—C9—C14	121.39 (18)
C6—C1—C2	120.9 (3)	C10—C9—C8	117.35 (19)
C6—C1—Cl1	120.46 (19)	C14—C9—C8	121.3 (2)
C2—C1—Cl1	118.6 (2)	C9—C10—C11	118.7 (2)
C3—C2—C1	120.1 (3)	C9—C10—H10	120.6
C3—C2—H2	120.0	C11—C10—H10	120.6
C1—C2—H2	120.0	O2—C11—C12	117.05 (18)
C2—C3—C4	120.2 (3)	O2—C11—C10	122.6 (2)
C2—C3—H3	119.9	C12—C11—C10	120.4 (2)
C4—C3—H3	119.9	C13—C12—C11	120.16 (18)
C3—C4—C5	120.4 (3)	C13—C12—H12	119.9

C3—C4—H4	119.8	C11—C12—H12	119.9
C5—C4—H4	119.8	O3—C13—C12	122.25 (18)
C4—C5—C6	121.0 (3)	O3—C13—C14	117.4 (2)
C4—C5—H5	119.5	C12—C13—C14	120.3 (2)
C6—C5—H5	119.5	C9—C14—C13	118.9 (2)
C5—C6—C1	117.4 (2)	C9—C14—H14	120.6
C5—C6—C7	121.1 (2)	C13—C14—H14	120.6
C1—C6—C7	121.5 (2)	H4A—O4—H4B	116.3
N1—C7—C6	120.9 (2)	H5A—O5—H5B	109.2

C7—N1—N2—C8	−172.3 (2)	O1—C8—C9—C10	−24.0 (3)
C6—C1—C2—C3	0.6 (4)	N2—C8—C9—C10	154.7 (2)
Cl1—C1—C2—C3	−178.9 (2)	O1—C8—C9—C14	156.3 (2)
C1—C2—C3—C4	−1.1 (5)	N2—C8—C9—C14	−25.0 (3)
C2—C3—C4—C5	0.6 (5)	C14—C9—C10—C11	2.8 (3)
C3—C4—C5—C6	0.4 (5)	C8—C9—C10—C11	−176.9 (2)
C4—C5—C6—C1	−0.9 (4)	C9—C10—C11—O2	178.0 (2)
C4—C5—C6—C7	−179.2 (3)	C9—C10—C11—C12	−1.0 (3)
C2—C1—C6—C5	0.4 (4)	O2—C11—C12—C13	179.4 (2)
Cl1—C1—C6—C5	179.9 (2)	C10—C11—C12—C13	−1.6 (4)
C2—C1—C6—C7	178.7 (2)	C11—C12—C13—O3	−177.8 (2)
Cl1—C1—C6—C7	−1.7 (3)	C11—C12—C13—C14	2.4 (4)
N2—N1—C7—C6	−179.62 (19)	C10—C9—C14—C13	−2.0 (4)
C5—C6—C7—N1	−3.9 (4)	C8—C9—C14—C13	177.7 (2)
C1—C6—C7—N1	177.8 (2)	O3—C13—C14—C9	179.6 (2)
N1—N2—C8—O1	−1.5 (4)	C12—C13—C14—C9	−0.6 (4)
N1—N2—C8—C9	179.82 (19)		

Hydrogen-bond geometry (Å, º)

D—H···*A*	*D*—H	H···*A*	*D*···*A*	*D*—H···*A*
O2—H2*B*···O5	0.82	1.87	2.674 (3)	168
O3—H3*A*···O1^{i}	0.82	1.85	2.665 (2)	169
N2—H2*A*···O2^{ii}	0.86	2.36	3.196 (3)	164
O4—H4*B*···O1	0.85	2.12	2.960 (4)	171
O5—H5*A*···O4^{iii}	0.85	1.99	2.816 (4)	163
O5—H5*B*···O3^{iv}	0.85	2.14	2.902 (3)	150

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