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Key indicators

Single-crystal X-ray study

$T = 294$ K

Mean $\sigma(\text{C}-\text{C}) = 0.003$ Å

R factor = 0.039

wR factor = 0.101

Data-to-parameter ratio = 15.6

For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.

N,N'-Bis(4-chlorobenzylidene)hydrazine

The title compound, $\text{C}_{14}\text{H}_{10}\text{Cl}_2\text{N}_2$, possesses a crystallographically imposed center of symmetry at the mid-point of the N—N bond and assumes a planar structure.

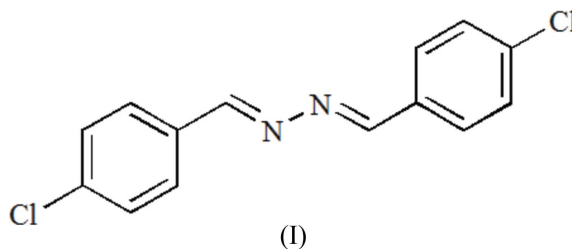
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Comment

Thus far a number of azine compounds containing both a diimine linkage and an N—N bond have been investigated in terms of their crystallography and coordination chemistry (Kundu *et al.*, 2005; Kesslen & Euler, 1999; Armstrong *et al.*, 1998; Xu *et al.*, 1997). We report here the crystal structure of the title compound, (I), in which the two 4-chlorobenzylidene units are directly linked, without a spacer unit, through the imine N atoms.



In the solid state, the molecule of (I) possesses a crystallographically imposed center of symmetry at the mid-point of the N—N bond. The C=N—N=C linkage is planar. The N—N bond [1.414 (3) Å] is slightly longer than in related azine compounds (Xu *et al.*, 2005; Liu *et al.*, 2004; Şengül *et al.*, 2004). The C=N—N angle [112.6 (2)°] is significantly smaller than the ideal sp^2 value of 120°, as a consequence of repulsion between the nitrogen lone pairs and the adjacent C=N bond. The whole molecule is essentially planar, the r.m.s deviation being 0.020 (2) Å.

Experimental

The title compound was synthesized by the reaction of 4-chlorobenzaldehyde with hydrazine hydrate in refluxing ethanol (Liu *et al.*, 2004). Single crystals suitable for X-ray analysis were obtained by slow evaporation of a cyclohexane solution.

Crystal data

$\text{C}_{14}\text{H}_{10}\text{Cl}_2\text{N}_2$

$M_r = 277.14$

Monoclinic, $P2_1/c$

$a = 3.9585$ (10) Å

$b = 6.9894$ (17) Å

$c = 23.150$ (6) Å

$\beta = 90.416$ (4)°

$V = 640.5$ (3) Å³

$Z = 2$

$D_x = 1.437$ Mg m⁻³

Mo $K\alpha$ radiation

Cell parameters from 1257

reflections

$\theta = 3.0$ – 25.3 °

$\mu = 0.49$ mm⁻¹

$T = 294$ (2) K

Block, colorless

$0.26 \times 0.20 \times 0.14$ mm

Data collection

Bruker SMART 1000 CCD area-detector diffractometer
 φ and ω scans
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.824$, $T_{\max} = 0.934$
 3387 measured reflections

1281 independent reflections
 944 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.025$
 $\theta_{\text{max}} = 26.3^\circ$
 $h = -3 \rightarrow 4$
 $k = -8 \rightarrow 7$
 $l = -28 \rightarrow 28$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.039$
 $wR(F^2) = 0.101$
 $S = 1.05$
 1281 reflections
 82 parameters
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0392P)^2 + 0.2929P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.001$
 $\Delta\rho_{\text{max}} = 0.32 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.31 \text{ e } \text{\AA}^{-3}$

Table 1

Selected geometric parameters (\AA , $^\circ$).

C1—C5	1.742 (2)	N1—N1 ⁱ	1.414 (3)
N1—C1	1.269 (3)		
C1—N1—N1 ⁱ	112.6 (2)	N1—C1—C2	121.8 (2)
N1 ⁱ —N1—C1—C2	−179.9 (2)	N1—C1—C2—C3	4.7 (3)
N1—C1—C2—C7	−176.1 (2)		

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

All H atoms were positioned geometrically and refined as riding, with C—H distances of 0.93 \AA and with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$.

Data collection: SMART (Bruker, 1997); cell refinement: SMART; data reduction: SAINT (Bruker, 1997); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Bruker, 1997); software used to prepare material for publication: SHELXTL.

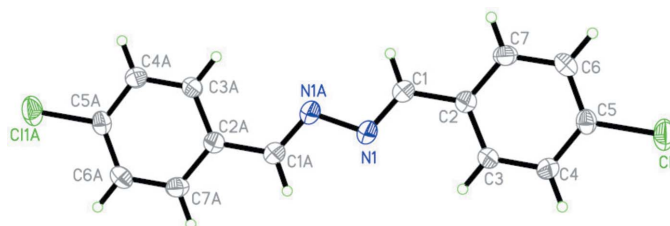


Figure 1

View of the molecule of (I), with displacement ellipsoids drawn at the 30% probability level. The suffix A corresponds to symmetry code (i) in Table 1.

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