Amide H atoms refined; others riding with fixed  $U_{\rm iso}$ 

Extinction correction: none Scattering factors from International Tables for Crystallography (Vol. C)

Table 1. Selected geometric parameters (Å, °)

		G12 G12	1 222 (2)
C1O1	1.225 (2)	C12—C13	1.377 (3)
C1—N2	1.337 (3)	C13—C14	1.360 (4)
C1-N1	1.390 (3)	C14—C15	1.353(4)
N1—C2	1.357 (3)	C15—C16	1.391 (3)
C2—O2	1.203 (2)	C21—C26	1.377 (3)
C2—C3	1.551 (3)	C21—C22	1.386 (3)
C3-N2	1.461 (3)	C22—C23	1.379 (3)
C3—C21	1.528 (3)	C23—N3	1.339 (3)
C3-C11	1.533 (3)	N3—C25	1.324 (3)
C11—C16	1.374 (3)	C25—C26	1.389 (3)
C11—C12	1.385 (3)		
O1C1N2	128.0 (2)	C16C11C3	120.9 (2)
01C1N1	124.1 (2)	C12C11C3	120.4 (2)
N2C1N1	107.8 (2)	C13C12C11	120.5 (2)
C2-N1-C1	112.4 (2)	C14—C13—C12	120.6 (3)
O2-C2-N1	127.6 (2)	C15-C14-C13	119.5 (2)
O2—C2—C3	126.2 (2)	C14C15C16	121.1 (3)
N1—C2—C3	106.1 (2)	C11C16C15	119.7 (2)
N2—C3—C21	112.7 (2)	C26C21C22	117.8 (2)
N2C3C11	110.75 (15)	C26C21C3	121.7 (2)
C21-C3-C11	112.9 (2)	C22C21C3	120.5 (2)
N2—C3—C2	100.7 (2)	C23C22C21	119.2 (2)
C21—C3—C2	108.08 (15)	N3—C23—C22	123.8 (2)
C11—C3—C2	111.0 (2)	C25—N3—C23	115.9 (2)
C1—N2—C3	112.7 (2)	N3—C25—C26	
C16C11C12	118.5 (2)	C21—C26—C25	118.4 (2)
210 211 -612	(2)	C2. C2., C2.,	(2)

Table 2. Hydrogen-bonding geometry (Å, °)

$D$ — $H \cdot \cdot \cdot A$	D—H	$\mathbf{H} \cdot \cdot \cdot \mathbf{A}$	$D \cdot \cdot \cdot A$	$D$ — $H \cdot \cdot \cdot A$
$N1-H1\cdots O1^{i}$	0.82(3)	2.02(3)	2.832 (2)	169 (3)
N2—H2···N3 <sup>ii</sup>	0.88 (3)	2.08 (3)	2.950(3)	173 (2)
Symmetry codes: (	i) $x - \frac{1}{2}, \frac{3}{2} - y$	z, $1-z$ ; (ii)	1-x,y-	$\frac{1}{2}, \frac{3}{2} - z$ .

Data collection: *P3/PC* (Siemens, 1989). Cell refinement: *P3/PC*. Data reduction: *SHELXTL* (Siemens, 1994). Program(s) used to solve structure: *SHELXS*86 (Sheldrick, 1990). Program(s) used to refine structure: *SHELXL*93 (Sheldrick, 1993). Molecular graphics: *SHELXTL*. Software used to prepare material for publication: *SHELXTL*.

Supplementary data for this paper are available from the IUCr electronic archives (Reference: BK1423). Services for accessing these data are described at the back of the journal.

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Acta Cryst. (1999). C55, 407-410

# 6-[*N*-(2-Hydroxyphenyl)aminomethylene]-cyclohexa-2,4-dien-1-one

ALOK K. MUKHERIEE, RAJIB LAL DE, INDRAJIT BANERJEE, CHITRA SAMANTA AND NIRMALYA P. NAYAK

<sup>a</sup>Department of Physics, Jadavpur University, Calcutta 700 032, India, and <sup>b</sup>Department of Chemistry, Jadavpur University, Calcutta 700 032, India. E-mail: akm@juphys.ernet.in

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### **Abstract**

The structure determination of the title compound,  $C_{13}H_{11}NO_2$ , establishes the tautomeric keto form of the salicylaldimine. The asymmetric unit consists of two crystallographically independent molecules which are essentially planar and are approximately orthogonal to each other. Strong intramolecular  $N-H\cdots O$  and intermolecular  $O-H\cdots O$  interactions influence the conformation of the molecules and the crystal packing. Intermolecular hydrogen bonds link the molecules in infinite chains.

## Comment

Salicylaldimines have been used extensively as ligands in coordination chemistry because of their diverse chelating ability (Long, 1995; Garnovski et al., 1993). The intramolecular hydrogen bond between O and N atoms in these systems plays a vital role in the formation of Schiff base compounds in the solid state by proton transfer from the hydroxyl-O atom to the imine-N atom (Hadjoudis et al., 1987; Elerman et al., 1997). The charge transfer through overlapping intermolecular  $\pi$ orbitals, with the associated change in the  $\pi$ -electron configuration in these compounds, also provides a basis for the development of molecular switches (Xu et al., 1994). This X-ray crystallographic study was undertaken in order to establish the solid-state molecular structure of (I) and to build up a hierarchy for N-substituted salicylaldimines.

 $C_{13}H_{11}NO_2$ 

A structure with this salicylaldimine in enol form was reported by Elerman et al. (1995), based on poor quality data (18% intensity decay and affected by disorder). In the present crystals, the salicylaldimine takes the tautomeric keto form. The asymmetric unit of the title compound, (I), consists of two crystallographically independent, but nearly identical, molecules (A and B) which are approximately orthogonal to each other: the dihedral angle between the essentially planar molecules A (r.m.s. deviation 0.117 Å) and B (r.m.s. deviation 0.120 Å) is 84.55 (3)°. The torsion angles C6—N1—C7—C8 179.4 (1) and C19—N2—C20—C21 179.9(1)° also illustrate that the molecules are almost planar. The C—O(hydroxyl) [1.348(2) and 1.354(2) Å in A and B, respectively] and N—C(phenyl) [1.412(2)] and 1.412(2) Å in A and B, respectively] bond lengths are consistent with typical values in related compounds (Elerman et al., 1995; Elmali & Elerman, 1997). The imine-N atoms (N1 and N2) are  $sp^2$  hybridized with bond angles summing to 360.0°. Each displays trigonal planar bonding geometry and the coplanarity of the resulting pseudo-rings (N1, C7, C8, C13, O2, H1N in A and N2, C20, C21, C26, O4, H2N in B) strongly suggests a resonance interaction in the pseudo-rings. This is reflected in the deviations of the C7—C8 [1.416(2)], C20-C21 [1.410(2)], C13-O2 [1.294(2)] and C26-O4 [1.297 (2) Å] bond lengths from standard C=C and C=O distances. A similar observation has been reported for 1-[N-(4-methyl-2-pyridyl)aminomethylene]-2(1H)-naphthalenone (Elerman et al., 1998).

Strong intra- and intermolecular N-H...O and O-H · O interactions influence the conformations of molecules A and B, and the crystal packing. The atoms H1N and H2N were located from a difference Fourier map at the end of the refinement process as small positive electron densities. In molecule A, the N1-H1N and H1N···O2 distances are 0.851 and 1.852 Å, respectively, while the corresponding values in B for N2—H2N and H2N···O4 are 0.867 and 1.850 Å, respectively. The planarity of the molecules in (I) makes it possible for the proton to transfer through the hydrogen bond with a small energy requirement (Bregman et al., 1964). Unlike the observations of Elerman et al. (1995) for the enol form of (I), N-(2-hydroxyphenyl)salicylaldimine, the present structure analysis does not show any disorder of the molecule or intramolecular O···O hydrogen bonding. The crystal packing is stabilized by intra- and intermolecular hydrogen bonds via oxo- and hydroxyl-O atoms. The molecules are linked by O—H···O hydrogen bonds into infinite chains with the sequence  $A \cdot \cdot \cdot B \cdot \cdot \cdot A \cdot \cdot \cdot B$ 

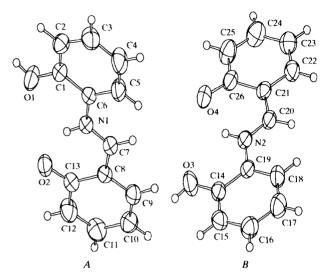


Fig. 1. ORTEPII (Johnson, 1976) view of the asymmetric unit of (I) with displacement ellipsoids shown at the 50% probability level.

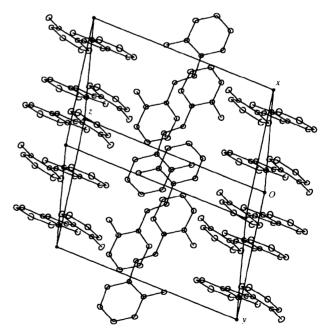


Fig. 2. The molecular packing in (I).

## **Experimental**

*N-p*-acetophenylsalicylaldimine (SACPNH) was prepared by condensation of equimolar quantities of salicylaldehyde and 4-aminoacetophenone in refluxing ethanol. Compound (I) was

obtained from the reaction of 2-aminophenol (0.002 mol, 0.22 g) with a solution of SACPNH (0.002 mol, 0.48 g) in 40 ml of ethanol. The orange-red precipitate was separated and recrystallized in methylene chloride-*n*-hexane solution.

## Crystal data

$C_{13}H_{11}NO_2$	Cu $K\alpha$ radiation
$M_r = 213.23$	$\lambda = 1.54180 \text{ Å}$
Triclinic	Cell parameters from 25
$P\overline{1}$	reflections
a = 9.034(2)  Å	$\theta = 4-75^{\circ}$
b = 10.155(1)  Å	$\mu = 0.758 \text{ mm}^{-1}$
c = 12.368 (2)  Å	T = 293 (2)  K
$\alpha = 110.35 (1)^{\circ}$	Prism
$\beta = 89.90 (2)^{\circ}$	$0.42 \times 0.34 \times 0.20 \text{ mm}$
$\gamma = 103.02 (1)^{\circ}$	Orange-red
$V = 1032.8 (3) \text{ Å}^3$	_
Z = 4	
$D_x = 1.371 \text{ Mg m}^{-3}$	

## Data collection

 $D_m$  not measured

Found Nonitor CAD 4	21.400
Enraf-Nonius CAD-4	3148 reflections with
diffractometer	$I > 2\sigma(I)$
$\omega$ –2 $\theta$ scans	$R_{\rm int} = 0.011$
Absorption correction:	$\theta_{\rm max} = 74.84^{\circ}$
empirical (North et al.,	$h = 0 \rightarrow 11$
1968)	$k = -12 \rightarrow 12$
$T_{\min} = 0.775, T_{\max} = 0.859$	$l = -15 \rightarrow 15$
3863 measured reflections	3 standard reflections
3676 independent reflections	every 100 reflections
-	intensity decay: <2%

# Refinement

Refinement on $F^2$	$\Delta \rho_{\text{max}} = 0.297 \text{ e Å}^{-3}$
$R[F^2 > 2\sigma(F^2)] = 0.050$	$\Delta \rho_{\min} = -0.294 \text{ e Å}^{-3}$
$wR(F^2) = 0.125$	Extinction correction:
S = 1.087	SHELXL93 (Sheldrick,
3676 reflections	1993)
312 parameters	Extinction coefficient:
H atoms: see text	0.0030 (7)
$w = 1/[\sigma^2(F_o^2) + (0.0948P)^2]$	Scattering factors from
where $P = (F_o^2 + 2F_c^2)/3$	International Tables for
$(\Delta/\sigma)_{\rm max} = 0.011$	Crystallography (Vol. C)

# Table 1. Selected geometric parameters (Å, °)

1.348(2)	C7—C8	1.416(2)
1.294(2)	C8—C9	1.415(2)
1.354(2)	C8—C13	1.432(2)
1.297(2)	C12—C13	1.415(2)
1.298(2)	C20—C21	1.410(2)
1.412(2)	C21—C22	1.410(2)
1.299(2)	C21—C26	1.438(2)
1.412(2)	C25—C26	1.413 (2)
127.06 (12)	C12C13C8	116.40 (14)
127.34 (12)	N2C20C21	123.33 (13)
122.99 (13)	C22C21C26	120.20 (14)
120.76 (13)	C25—C26—C21	116.51 (15)
-2.5(2)	O3C14C19N2	1.7(2)
179.4(1)	C19-N2-C20-C21	179.9(1)
-1.0(2)	N2C20C21C26	0.8(2)
-3.5(2)	C20—C21—C26—O4	3.4(2)
	1.294 (2) 1.354 (2) 1.297 (2) 1.298 (2) 1.412 (2) 1.299 (2) 1.412 (2) 127.06 (12) 127.34 (12) 120.76 (13) -2.5 (2) 179.4 (1) -1.0 (2)	1.294 (2)

Table 2. Hydrogen-bonding geometry (Å, °)

$D$ — $H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	$D$ — $H \cdot \cdot \cdot A$
NI-HIN···O2	2.597(2)	145.1(1)
N2H2N· · · O4	2.595(2)	143.0(1)
O1H1OO4 <sup>i</sup>	2.585(2)	161.1(1)
O3—H3O· · · O2 <sup>n</sup>	2.571(2)	162.2(1)

Symmetry codes: (i) 1 - x, -y, 2 - z; (ii) 2 - x, -y, 2 - z.

The structure was solved by direct methods and refined by full-matrix anisotropic least squares without any constraint. H atoms were located from the difference Fourier maps; their positions were then idealized and treated as riding, and their isotropic displacement parameters were refined.

Data collection: CAD-4 Software (Enraf-Nonius, 1989). Cell refinement: CAD-4 Software. Data reduction: CAD-4 Software. Program(s) used to solve structure: MULTAN88 (Debaerdemaeker et al., 1988). Program(s) used to refine structure: SHELXL93 (Sheldrick, 1993). Molecular graphics: ZORTEP (Zsolnai & Pritzkow, 1995). Software used to prepare material for publication: SHELXL93.

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: HA1241). Services for accessing these data are described at the back of the journal.

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Acta Cryst. (1999). C55, 410-411

# 1,4-Diformyl-*t*-5-methyl-*r*-2,*t*-3-diphenyl-piperazine-1,4-dicarbaldehyde

P. LAAVANYA, K. PANCHANATHESWARAN, R. MURUGADOSS AND R. JEYARAMAN

Department of Chemistry, Bharathidasan University, Tiruchirappalli 620 024, India. E-mail: pan@bdu.ernet.in

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#### **Abstract**

A single-crystal X-ray diffraction study of the title compound  $(C_{19}H_{20}N_2O_2)$  confirms the chair conformation of the piperazine ring, with the methyl and phenyl groups in axial orientations.

### Comment

The major highlight of this investigation is the presence of an alternate chair conformation of the piperazine ring of the title compound, (I), in which the phenyl and the methyl groups assume axial orientations. The nor-

mal chair conformation with equatorial substituents is not preferred due to the steric interaction between the N-formyl and phenyl groups. The C5—C6 and C3—C2 bond distances, 1.519 (3) and 1.537 (3) Å, respectively, are in the range 1.515–1.535 Å reported for other piperazine derivatives (Sekido et al., 1985; Okamato et al., 1979, 1982; Sakurai et al., 1977; Davis & Hassel, 1963). The bond angles around the  $Csp^3$  atoms in the piperazine ring  $[109.0 (2)-111.3 (2)^\circ]$  indicate only slight distortion from the tetrahedral arrangement of groups around these atoms. The bond angles around N1 and N4  $[116.6 (2)-122.2 (2)^\circ]$  suggest  $sp^2$  character for these atoms, as in the case of similar piperazine derivatives (Sekido et al., 1985; Okamato et al., 1979; Sakurai et al., 1977). The N1 and N4 atoms are at a dis-

tance of 0.554 (2) and -0.594 (2) Å, respectively, from the C2—C3—C5—C6 plane, indicating a chair conformation in which the two N atoms are unsymmetrically disposed. The axial orientations of the methyl and the two phenyl groups are evident from the torsion angles, N4—C5—C6—C9 76.9 (2), N1—C2—C3—C16—78.0 (2) and N4—C3—C2—C10—78.2 (2)°.

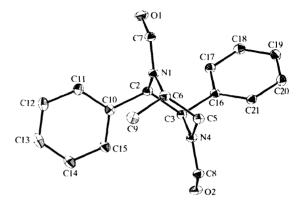


Fig. 1. ZORTEP (Zsolnai, 1997) plot of (I) showing 30% probability displacement ellipsoids. The H atoms are omitted for clarity.

# **Experimental**

The title compound was obtained by the formylation of t-5-methyl-r-2,t-3-diphenylpiperazine by a mixture of acetic anhydride and 85% formic acid using benzene as solvent. The crude product isolated from the benzene layer was recrystallized using ethanol.

# Crystal data

$C_{19}H_{20}N_2O_2$	Cu $K\alpha$ radiation
$M_r = 308.37$	$\lambda = 1.54180 \text{ Å}$
Monoclinic	Cell parameters from 25
$P2_1/n$	reflections
a = 12.355(3)  Å	$\theta = 4.12-70.13^{\circ}$
b = 11.645(3)  Å	$\mu = 0.657 \text{ mm}^{-1}$
c = 12.750(3)  Å	T = 293 (2)  K
$\beta = 117.407 (16)^{\circ}$	Cubic
$V = 1628.5 (7) \text{ Å}^3$	$0.1 \times 0.1 \times 0.1 \text{ mm}$
Z = 4	Colourless
$D_x = 1.258 \text{ Mg m}^{-3}$	
$D_m$ not measured	

# Data collection

Rigaku AFC-7S diffractom-	2382 reflections with
eter	$I > 2\sigma(I)$
$\omega$ –2 $\theta$ scans	$R_{\rm int} = 0.060$
Absorption correction:	$\theta_{\rm max} = 70.13^{\circ}$
$\psi$ scans (North et al.,	$h=0 \rightarrow 15$
1968)	$k = 0 \rightarrow 14$
$T_{\text{min}} = 0.829, T_{\text{max}} = 0.936$	$l = -15 \rightarrow 13$
3358 measured reflections	3 standard reflections
3021 independent reflections	every 150 reflections
•	intensity decay: negligible