# organic papers

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#### Key indicators

Single-crystal X-ray study T = 296 KMean  $\sigma$ (C–C) = 0.003 Å R factor = 0.038 wR factor = 0.106 Data-to-parameter ratio = 14.1

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

© 2002 International Union of Crystallography Printed in Great Britain – all rights reserved (Z)-4-(4-{4-[(Z)-1-Methyl-3-oxobut-1-enylamino]phenoxy}phenylamino)pent-3-en-2-one

In the structure of the title compound,  $C_{22}H_{24}N_2O_3$ , short intermolecular  $C-H\cdots O$  contacts influence the crystal packing and molecular ladders are formed. Intramolecular  $N-H\cdots O$  hydrogen bonds make the Z configuration stable by formation of six-membered rings.

#### Comment

Schiff bases are becoming increasingly important as biological, analytical, pharmacological and antimicrobial reagents in coordination chemistry. There are many reports of Schiff bases obtained by condensation between the  $NH_2$  group of an amine and the carbonyl group of aldehydes/ketones (Ismail, 2000; Raman *et al.*, 2001; Daniel Thangadurai & Natarajan, 2000). A literature survey revealed fewer results involving isomers of Schiff bases. We obtained the title compound, (I), by the usual type of condensation; however, the expected C-C=N group is, in fact, C=C-N.



A view of the molecule of (I) is shown in Fig. 1. The dihedral angle between the two planes defined by O1 and each of the two phenyl rings is 73.39 (7)°. The N1–H1N···O2 and N2–H2N···O3 hydrogen bonds stabilize the Z configuration of the C=C bonds by forming six-membered rings (Table 2). The torsion angles C7–C9–C10–O2 [1.3 (3)°] and C7–C9–C10–C11 [–177.81 (19)°] indicate that the group N1/C7/C8/C9/C10/C11/O2 is nearly planar; the group N2/C18/C19/C20/C21/C22/O3 is slightly more non-planar, as is evident from the torsion angles C18–C20–C21–O3 [–2.3 (3)°] and C18–C20–C21–C22 [175.70 (18)°]. The dihedral angle between these two groups is 88.21 (6)°. The two halves of the molecule resemble each other in geometry (Table 1).

The short intermolecular  $C-H\cdots O$  contacts (Table 2) involving both ketone functions are respectively 0.19, 0.18 and 0.12 Å shorter than the sum of van der Waals radii (Bondi, 1964). They may be classified as weak to very weak hydrogen bonds (Desiraju & Steiner, 1999), and the second is near-linear (Jeffrey *et al.*, 1985). Examination of the structure with *PLATON* (Spek, 2002) shows that there are no solvent-accessible voids.

The intermolecular  $C-H\cdots O$  contacts link molecules into dimers, dimers face-to-face to form four-molecule aggregates, and aggregates into ladders, to give a two-dimensional network (Fig. 2).

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#### Figure 1

A view of the molecular structure of (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level and H atoms are shown as spheres of arbitrary radii.

#### **Experimental**

The title compound was prepared by the condensation in ethanol of 4,4'-diamino-diphenyl ether (0.5 mol) with pentane-2,4-dione (1.0 mol). The solution was stirred for 2 h at 433 K. Crystals suitable for diffraction study were obtained after one month (m.p. 428 K). Spectroscopic analysis, <sup>1</sup>H NMR (CDCl<sub>3</sub>,  $\delta$ , p.p.m.): 1.97 (*s*, 6H), 2.10 (*s*, 6H), 5.19 (*s*, 1H), 7.04 (*m*, 8H), 12.39 (*s*, 1H).

#### Crystal data

$C_{22}H_{24}N_2O_3$ $M_r = 364.43$ Monoclinic, $P2_1/c$ a = 8.769 (1) Å b = 20.073 (4) Å c = 11.928 (2) Å $\theta_{-} = 107.52$ (1)°	$D_x = 1.209 \text{ Mg m}^{-3}$ Mo $\kappa \alpha$ radiation Cell parameters from 26 reflections $\theta = 3.2-15.2^{\circ}$ $\mu = 0.08 \text{ mm}^{-1}$ T = 206 (2) K
p = 107.55(1)	1 = 250(2) K
V = 2002.1 (6) A <sup>2</sup>	Block, yellow
Z = 4	$0.54 \times 0.52 \times 0.44$ mm
Data collection	
Siemens P4 diffractometer	$\theta_{\rm max} = 25.0^{\circ}$
(i) scans	$h = 0 \rightarrow 10$
Absorption correction: none	$k = 0 \rightarrow 10$ $k = 0 \rightarrow 23$
3762 massured reflections	$k = 0 \rightarrow 25$ $l = -14 \rightarrow 13$
3520 independent reflections	$i = -14 \rightarrow 15$
2220 independent reflections	5 standard reflections
2558 reflections with $I > 2\sigma(I)$	every 97 renections
$R_{\rm int} = 0.010$	intensity decay: 4.4%
Refinement	
Refinement on $F^2$	$w = 1/[\sigma^2(F_o^2) + (0.0609P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.038$	where $P = (F_0^2 + 2F_c^2)/3$
$wR(F^2) = 0.106$	$(\Delta/\sigma)_{\rm max} < 0.001$
S = 0.98	$\Delta \rho_{\rm max} = 0.16  {\rm e}  {\rm \AA}^{-3}$
3520 reflections	$\Delta \rho_{\rm min} = -0.15  {\rm e}  {\rm \AA}^{-3}$
249 parameters	Extinction correction: SHELXL97
H-atom parameters constrained	Extinction coefficient: 0.0223 (19)
ri atom parameters constrained	Extinction coefficient. $0.0223$ (19)





### Table 1

O1-C1	1.3911 (19)	N1-C4	1.417 (2)
O1-C12	1.3926 (19)	N2-C18	1.335 (2)
O2-C10	1.245 (2)	N2-C15	1.419 (2)
O3-C21	1.2406 (19)	C7-C9	1.368 (2)
N1-C7	1.339 (2)	C18-C20	1.373 (2)
C1 - O1 - C12	117.75 (12)	N1-C7-C8	119.99 (15)
C7-N1-C4	130.90 (14)	C14-C15-N2	123.45 (15)
C18-N2-C15	131.37 (14)	C16-C15-N2	117.30 (15)
C5-C4-N1	122.85 (15)	N2-C18-C20	119.66 (15)
C3-C4-N1	118.10 (14)	N2-C18-C19	119.41 (16)
N1-C7-C9	119.59 (15)		
C4-N1-C7-C9	175.96 (17)	C15-N2-C18-C20	-178.18 (16)
N1-C7-C9-C10	-3.6(3)	N2-C18-C20-C21	3.3 (3)
C7-C9-C10-O2	1.3 (3)	C18-C20-C21-O3	-2.3(3)
C7-C9-C10-C11	-177.81 (19)	C18-C20-C21-C22	175.70 (18)

#### Table 2

Contact distances (Å).

H8C···O2 <sup>i</sup>	2.54	$H16 \cdots O2^{ii}$	2.60
Symmetry codes: (i) $x, \frac{1}{2} - y, z - y$	$\frac{1}{2}$ ; (ii) x -	$1, \frac{1}{2} - y, z - \frac{1}{2}$	

# Table 3

Hydrogen-bonding geometry (Å,  $^\circ).$ 

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
N1-H1N···O2	0.86	1.94	2.643 (2)	139
$N2 - H2N \cdot \cdot \cdot O3$	0.86	1.92	2.637 (2)	139
$C6-H6\cdots O3^{i}$	0.93	2.53	3.175 (2)	127

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

All H atoms were generated geometrically and refined using a riding model, with C-H = 0.93 Å for aromatic H atoms and 0.96 Å for methyl H atoms, N-H = 0.86 Å, and  $U_{\rm iso}({\rm H}) = 1.2U_{\rm eq}$  of the parent atoms.

Data collection: *XSCANS* (Fait, 1991); cell refinement: *XSCANS*; data reduction: *SHELXTL* (Sheldrick, 1997*b*); program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1990); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997*a*); molecular graphics: *ORTEP*-3 for Windows (Farrugia, 1997); software used to prepare material for publication: *SHELXTL*.

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