Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

## Shi Jun‡

Department of Basic Course, Tianjin Agricultural College, Tianjin 300384, People's Republic of China

**‡** Current address: School of Chemical Engineering and Technology, Tianjin University, Tianjin 300072, People's Republic of China.

Correspondence e-mail: shi\_jun99@163.com

#### Key indicators

Single-crystal X-ray study T = 294 KMean  $\sigma(C-C) = 0.003 \text{ Å}$  R factor = 0.050 wR factor = 0.127 Data-to-parameter ratio = 15.5

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

# (*E*)-1-[4-(Benzyloxy)benzylidene]-2-(4-nitrophenyl)hydrazine

The title compound,  $C_{20}H_{17}N_3O_3$ , was prepared by the reaction of 4-(benzyloxy)benzaldehyde and 1-(4-nitrophenyl)hydrazine. In the crystal structure, the central benzene ring makes dihedral angles of 3.16 (12) and 64.87 (6)° with the *p*-nitrophenyl and phenyl rings, respectively. The crystal packing is stabilized by intermolecular N-H···O hydrogen bonds. Received 10 October 2005 Accepted 24 October 2005 Online 27 October 2005

### Comment

In continuation of our search for new Schiff bases functioning as ligands (Jun, 2005), we present the crystal structure of the title compound, (I).



In (I) (Fig. 1), the bond lengths and angles (Table 1) are normal. The benzene ring (C8–C13) makes dihedral angles of 64.87 (6) and 3.16 (12)° with the phenyl (C1–C6) and benzene (C15–C20) rings, respectively. The crystal packing (Fig. 2) is stabilized by the intermolecular  $N-H\cdots$ O hydrogen bonds (Table 2).

# **Experimental**

An anhydrous ethanol solution of 4-(benzyloxy)benzaldehyde (2.12 g,10 mmol) was added to an anhydrous ethanol solution of 1-(4nitrophenyl)hydrazine (1.53 g, 10 mmol), and the mixture was stirred at 350 K for 5 h under nitrogen. A yellow precipitate was isolated, recrystallized from ethanol, and then dried *in vacuo* to give the pure compound in 87% yield. Yellow single crystals suitable for X-ray analysis were obtained by slow evaporation of an ethanol solution of (I).



**Figure 1** The structure of (I), with displacement ellipsoids for non-H atoms drawn at the 30% probability level.

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# organic papers

#### Crystal data

 $\begin{array}{l} C_{20}H_{17}N_{3}O_{3}\\ M_{r}=347.37\\ \text{Monoclinic, }P2_{1}/c\\ a=6.148\ (3) \text{ Å}\\ b=36.651\ (15) \text{ Å}\\ c=7.938\ (3) \text{ Å}\\ \beta=93.551\ (7)^{\circ}\\ V=1785.2\ (13) \text{ Å}^{3}\\ Z=4 \end{array}$ 

### Data collection

Bruker SMART CCD area-detector
diffractometer
$\varphi$ and $\omega$ scans
Absorption correction: multi-scan
(SADABS; Sheldrick, 1996)
$T_{\min} = 0.958, T_{\max} = 0.982$
10028 measured reflections

#### Refinement

<b>P</b> 2 <b>P</b> <sup>2</sup>	
Refinement on F <sup>2</sup>	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.050$	$w = 1/[\sigma^2(F_o^2) + (0.0516P)^2]$
$wR(F^2) = 0.127$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.00	$(\Delta/\sigma)_{\rm max} = 0.001$
3643 reflections	$\Delta \rho_{\rm max} = 0.14 \ {\rm e} \ {\rm \AA}^{-3}$
235 parameters	$\Delta \rho_{\rm min} = -0.17 \text{ e} \text{ \AA}^{-3}$

 $D_x = 1.292 \text{ Mg m}^{-3}$ 

Cell parameters from 1822

 $0.40 \times 0.34 \times 0.20 \text{ mm}$ 

3643 independent reflections 1738 reflections with  $I > 2\sigma(I)$ 

Mo  $K\alpha$  radiation

reflections  $\theta = 2.6-22.7^{\circ}$ 

 $\mu = 0.09 \text{ mm}^{-1}$ 

T = 294 (2) K

Block, yellow

 $\begin{aligned} R_{\text{int}} &= 0.051\\ \theta_{\text{max}} &= 26.5^{\circ}\\ h &= -7 \rightarrow 7\\ k &= -18 \rightarrow 45\\ l &= -9 \rightarrow 9 \end{aligned}$ 

# Table 1

Selected geometric parameters (Å, °).

O1-C8	1.374 (2)	N1-C14	1.284 (2)
D1-C7 1.433 (2)		N1-N2	1.368 (2)
O2-N3	1.238 (3)	N2-C15	1.368 (3)
O3-N3	1.227 (3)	N3-C18	1.453 (3)
C8-O1-C7	117.79 (16)	O3-N3-O2	123.5 (3)
C14-N1-N2	117.18 (19)	O3-N3-C18	119.1 (3)
C15-N2-N1	120.48 (19)	O2-N3-C18	117.4 (3)

### Table 2

Hydrogen-bond geometry (Å,  $^{\circ}$ ).

$N2-H2\cdots O2^{i}$ 0.86 2.06 2.912 (3) 170	$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots \mathbf{A}$
	$N2-H2\cdots O2^{i}$	0.86	2.06	2.912 (3)	170

Symmetry code: (i)  $x - 1, -y + \frac{1}{2}, z + \frac{1}{2}$ .

# Figure 2

View of the crystal packing, with the intermolecular  $N\!-\!H\!\cdots\!O$  hydrogen bonds shown by dashed lines.

All H atoms were included in calculated positions and refined using a riding-model approximation, with C–H = 0.93–0.97 Å, N–H = 0.86 Å and  $U_{\rm iso}({\rm H}) = 1.2 U_{\rm eq}({\rm C,N})$ .

Data collection: *SMART* (Bruker, 1999); cell refinement: *SAINT* (Bruker, 1999); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997*a*); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997*a*); molecular graphics: *SHELXTL* (Sheldrick, 1997*b*); software used to prepare material for publication: *SHELXTL*.

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