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# K. R. Krishnapriya,<sup>a</sup> N. Sampath,<sup>b</sup> S. Aravindhan,<sup>b</sup> M. N. Ponnuswamy<sup>b</sup>\* and M. Kandaswamy<sup>a</sup>

<sup>a</sup>Department of Inorganic Chemistry, University of Madras, Guindy Campus, Chennai 600025, India, and <sup>b</sup>Department of Crystallography and Biophysics, University of Madras, Guindy Campus, Chennai 600025, India

Correspondence e-mail: mnpsy2004@yahoo.com

#### **Key indicators**

Single-crystal X-ray study T = 293 K Mean  $\sigma$ (C–C) = 0.006 Å R factor = 0.059 wR factor = 0.198 Data-to-parameter ratio = 13.5

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

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# 2,4-Bis(pyridine-2-carbonyl)-1-(2-pyridyl)-3,5-di-p-tolylcyclohex-1-ol

In the title compound,  $C_{37}H_{33}N_3O_3$ , the central cyclohexane ring adopts a chair conformation. The molecular structure is stabilized by C-H···N, C-H···O, O-H···N and  $\pi$ - $\pi$ interactions. Interestingly,  $\pi$ - $\pi$  and C-H··· $\pi$  weak interactions play a major role in the crystal packing. Received 14 October 2004 Accepted 8 November 2004 Online 20 November 2004

#### Comment

The crystal structure determination of the title compound, (I), was carried out in order to elucidate the molecular conformation and hydrogen-bonding characteristics.



The cyclohexane ring adopts a chair conformation, with puckering parameters (Cremer & Pople, 1975) Q =0.567 (4) Å,  $\theta = 1.8$  (4)° and  $\varphi = 353$  (8)°. The pyridine-2carbonyl group attached to C33 and the hydroxy group are in axial orientations, whereas the other substituents are attached to the cyclohexane ring in equatorial positions (Fig. 1). The C12-C17 benzene ring makes dihedral angles of 35.3 (2) and 75.1 (2) $^{\circ}$ , respectively, with the N1/C18–C22 and N2/C6–C10 pyridine rings. The C24-C29 benzene ring makes dihedral angles of 83.8 (2) and 51.6 (2) $^{\circ}$ , respectively, with the N1/C18– C22 and N3/C1-C5 pyridine rings. The dihedral angle between the N2/C6-C10 and N3/C1-C5 pyridine rings is 21.5 (2)°. In the cyclohexane ring, the C30-C31 [1.564 (6) Å] and C32-C33 [1.568 (5) Å] bond distances are longer than normal, due to the electron-acceptor capability of the acetylpyridine groups and strain induced by bulky substituents on the cyclohexane ring (Thomson et al., 1995; Vasilyev et al., 1990)





A ZORTEP (Zsolnai, 1998) plot of (I), showing 50% probability displacement ellipsoids and the atomic numbering.



#### Figure 2

Packing of the molecules viewed down the *a* axis. The dashed lines indicate the  $\pi$ - $\pi$  and C-H··· $\pi$  interactions.

The molecular structure is stabilized by intramolecular C- $H \cdots N$ ,  $C - H \cdots O$  and  $O - H \cdots N$  hydrogen bonds (Table 2) and a  $\pi$ - $\pi$  interaction between the N2/C6-C10 and N3/C1-C5 pyridine rings [centroid–centroid distance = 3.657 (4) Å]. The  $\pi$ - $\pi$  and weak C-H··· $\pi$  interactions (Desiraju, 1989) play a major role in the crystal packing (Fig. 2), in addition to van der Waals forces. The N3/C1-C5 and N2/C6-C10 pyridine rings of the glide-related molecule at  $(\frac{1}{2} + x, \frac{1}{2} - y, z - \frac{1}{2})$  are stacked with a centroid-centroid separation of 3.898 (4) Å. A C- $H \cdots \pi$  interaction,  $C4 - H4 \cdots Cg1^i$  (Cg1 is the C12-C17 ring centroid), is also observed between these two molecules (Table 2). The weak interactions result in the formation of a zigzag chain along the c axis (Fig. 2). In the chain, molecules translated by a unit along the c axis are linked via C20-H20···Cg2<sup>ii</sup> (Cg2 is the C24–C29 ring centroid) interactions. Adjacent chains are interlinked through C36-H36C···Cg2<sup>iii</sup> interactions. All symmetry codes are as given in Table 2.

# **Experimental**

p-Tolualdehyde (0.72 g, 5.99 mmol) in ethanol (30 ml) was added dropwise to a stirred solution of NaOH (0.6 g) in water (6 ml). 2-Acetylpyridine (1.089 g, 8.99 mmol) was added dropwise and the solution was stirred at room temperature for 80 h. The precipitate formed during this period was filtered off and washed with water and then with diethyl ether. The crude product was recrystallized from chloroform to give colourless crystals.

#### Crystal data

C <sub>37</sub> H <sub>33</sub> N <sub>3</sub> O <sub>3</sub>	$D_x = 1.255 \text{ Mg m}^{-3}$
$M_r = 567.66$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/n$	Cell parameters from 25
a = 8.534 (7)  Å	reflections
b = 34.067 (10)  Å	$\theta = 2.1 - 25.0^{\circ}$
c = 10.458 (5)  Å	$\mu = 0.08 \text{ mm}^{-1}$
$\beta = 98.91 \ (5)^{\circ}$	T = 293 (2) K
$V = 3004 (3) \text{ Å}^3$	Needle, colourless
Z = 4	$0.41\times0.15\times0.12~\text{mm}$

# Data collection

 $\theta_{\rm max} = 25.0^{\circ}$ Enraf-Nonius CAD-4  $h=0\rightarrow 10$ diffractometer  $\omega$  scans  $k = 0 \rightarrow 40$  $l = -12 \rightarrow 12$ Absorption correction: none 5638 measured reflections 3 standard reflections 5267 independent reflections 2164 reflections with  $I > 2\sigma(I)$  $R_{\rm int} = 0.017$ 

### Refinement

Refinement on  $F^2$  $R[F^2 > 2\sigma(F^2)] = 0.059$  $wR(F^2) = 0.198$ S = 0.945267 reflections 391 parameters

## Table 1

Selected geometric parameters (Å, °).

C1-N3	1.338 (5)	C18-N1	1.338 (5)
C5-N3	1.328 (6)	C22-N1	1.331 (5)
C6-N2	1.334 (5)	C23-O2	1.211 (5)
C6-C11	1.499 (6)	C30-O3	1.423 (4)
C11-O1	1.221 (5)		
N3-C1-C30	115.3 (4)	C13-C12-C32	123.3 (4)
C2-C1-C30	123.9 (4)	N1-C18-C19	123.4 (4)
N3-C5-C4	124.8 (5)	N1-C18-C23	117.7 (4)
N2-C6-C7	122.6 (4)	N1-C22-C21	124.1 (5)
N2-C6-C11	116.3 (4)	O2-C23-C33	122.7 (4)
N2-C10-C9	124.5 (5)	C18-C23-C33	118.4 (4)
O1-C11-C31	122.5 (4)	C29-C24-C25	116.4 (4)
C6-C11-C31	117.7 (4)	C29-C24-C34	123.8 (4)
C17-C12-C13	116.6 (4)		
C35-C30-C31-C32	-56.4(4)	C32-C33-C34-C35	52.7 (4)
C30-C31-C32-C33	54.7 (4)	C33-C34-C35-C30	-56.4(4)
C31-C32-C33-C34	-52.4 (4)	C31-C30-C35-C34	56.7 (4)

every 100 reflections

intensity decay: none

H-atom parameters constrained

 $w = 1/[\sigma^2(F_o^2) + (0.1021P)^2]$ 

 $(\Delta/\sigma)_{\rm max} = 0.001$ 

 $\Delta \rho_{\rm max} = 0.23 \ {\rm e} \ {\rm \AA}^2$ 

 $\Delta \rho_{\rm min} = -0.24 \text{ e } \text{\AA}^{-3}$ 

where  $P = (F_o^2 + 2F_c^2)/3$ 

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Table 2 Hydrogen-bonding geometry (Å,  $^\circ).$ 

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
O3−H3A…N3	0.82	2.10	2.586 (5)	118
C31-H31···N2	0.98	2.43	2.887 (5)	108
C33-H33···N1	0.98	2.38	2.890 (6)	112
C35−H35 <i>B</i> ···O2	0.97	2.46	3.070 (6)	121
$C4-H4\cdots Cg1^{i}$	0.93	3.04	3.805 (6)	141
C20−H20···Cg2 <sup>ii</sup>	0.93	3.04	3.824 (6)	143
$C36-H36C\cdots Cg2^{iii}$	0.96	2.94	3.685 (6)	135

Symmetry codes: (i)  $\frac{1}{2}$  + x,  $\frac{1}{2}$  - y, z -  $\frac{1}{2}$ ; (ii) x, y, 1 + z; (iii) 2 - x, -y, 1 - z. Cg1 is the C12-C17 ring centroid and Cg2 is the C24-C29 ring centroid

The high proportion of unobserved data may be a result of the high mosaicity of the crystal. All the H atoms were positioned geometrically (O-H = 0.82 Å and C-H = 0.93–0.98 Å) and allowed to ride on their parent atoms, with  $U_{\rm iso}(\rm H) = 1.2$  or 1.5 times  $U_{\rm eq}(\rm parent atom)$ .

Data collection: CAD-4 Software (Enraf-Nonius, 1989); cell refinement: CAD-4 Software; data reduction: XCAD4 (Harms & Wocadlo, 1995); program(s) used to solve structure: SHELXS97

(Sheldrick, 1997); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *ZORTEP* (Zsolnai, 1998) and *PLATON* (Spek, 2003); software used to prepare material for publication: *SHELXL*97 and *PARST* (Nardelli, 1995).

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