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Pyridine-4-carbaldehyde azine

S. Shanmuga Sundara Raj,^a Hoong-Kun Fun,^{a*} Jing Zhang,^b Ren-Geng Xiong^b and Xiao-Zeng You^b

^aX-ray Crystallography Unit, School of Physics, Universiti Sains Malaysia, 11800 USM, Penang, Malaysia, and ^bCoordination Chemistry Institute & State Key Laboratory of Coordination Chemistry, Nanjing University, Nanjing 210093, People's Republic of China

Correspondence e-mail: hkfun@usm.my

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In the crystal structure of the title compound, $C_{12}H_{10}N_4$, the pyridine ring makes a dihedral angle of $1.12~(9)^\circ$ with the mean plane of the complete almost planar and crystal-lographically centrosymmetric molecule. There are stacks of parallel molecules along the *a*-axis direction, with alternate stacks having a herring-bone arrangement relative to each other and an interplanar spacing of 3.551~Å.

Comment

The title compound, (I), is known not only as a coordinating reagent to determine the concentration of the Fe^{II} ion in solution (Luque de Castro & Valcarcel, 1978), but also as a precursor in the atactic polymerization of 4-vinylpyridine (Biedermann et al., 1972). In addition, this Schiff base ligand derived from hydrazine finds a wide range of applications in coordination chemistry owing to its polydentate chelating ability (Tarafder & Khan, 1991). Although the synthesis and characterization of 4,4-azinodimethyldipyridine and the crystal structure of its diperchlorate salt, $C_{12}H_{10}N_4^{2+} \cdot 2ClO_4^{-}$, have been reported (Chen et al., 1997), the crystal structure of 4,4'-azinodimethyldipyridine or pyridine-4-carbaldehyde azine, (I), has not been reported. The title compound acts as a potential bridging ligand in the study of supramolecular chemistry and can coordinate to metal ions to form multidimensional structures.

There is half a molecule in the asymmetric unit and the other half is inversion related. The bond distances and angles agree with those of the perchlorate salt (Chen *et al.*, 1997). The C6—N2 distance is shorter than 1.285 (7) Å and the bond

angles C6–N2–N2ⁱ [symmetry code: (i) 1-x, 2-y, 1-z] and C1–C6–N2 are slightly smaller than those of 112.2 (2) and 120.7 (2)°, respectively, reported for 2,2′-azinodimethyl-diphenol (Xu *et al.*, 1994). The molecule adopts an antiperiplanar conformation with respect to the N2–N2ⁱ bond. Atoms C6 and C6ⁱ are in a *trans* orientation with respect to the N2–N2ⁱ bond. The pyridine ring makes a dihedral angle of 1.12 (9)° with the mean plane of the complete almost planar molecule. There are stacks of parallel molecules along the *a*-axis direction, with alternate stacks having a herring-bone arrangement relative to each other with an interplanar spacing of 3.551 Å.

Experimental

The title compound was prepared by stirring hydrazine hydrate (80%) and pyridine-4-carboxaldehyde under flowing N_2 in anhydrous EtOH solution at 353 K for 6 h. The solution was then cooled and filtered. The filtrate was evaporated at room temperature for a few weeks and orange block-shaped crystals were obtained.

Crystal data

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$C_{12}H_{10}N_4$	$D_x = 1.293 \text{ Mg m}^{-3}$
$M_r = 210.24$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/c$	Cell parameters from 2605
a = 3.8515 (3) Å	reflections
b = 11.0201 (8) Å	$\theta = 2.45 - 28.32^{\circ}$
c = 12.7303 (9) Å	$\mu = 0.083 \text{ mm}^{-1}$
$\beta = 92.312 (2)^{\circ}$	T = 293 (2) K
$V = 539.88 (7) \text{ Å}^3$	Block, orange
Z = 2	$0.38 \times 0.36 \times 0.32 \text{ mm}$

Data collection

Siemens SMART CCD area-	$R_{\rm int} = 0.031$
detector diffractometer	$\theta_{\rm max} = 28.42^{\circ}$
ω scans	$h = -5 \rightarrow 5$
3584 measured reflections	$k = -14 \rightarrow 7$
1325 independent reflections	$l = -16 \rightarrow 16$
1065 reflections with $I > 2\sigma(I)$	

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0958P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.067$	+ 0.1438P]
$wR(F^2) = 0.110$	where $P = (F_o^2 + 2F_c^2)/3$
S = 1.247	$(\Delta/\sigma)_{\rm max} < 0.001$
1325 reflections	$\Delta \rho_{\text{max}} = 0.31 \text{ e Å}^{-3}$
93 parameters	$\Delta \rho_{\min} = -0.20 \text{ e Å}^{-3}$
All H-atom parameters refined	

Table 1 Selected geometric parameters (Å, °).

1.332 (3)	N2-C6	1.266 (3)
1.337 (3)	$N2-N2^{i}$	1.407 (3)
116.7 (2)	C6-N2-N2'	112.2 (2)
		1.337 (3) N2-N2 ⁱ

Symmetry code: (i) 1 - x, 2 - y, 1 - z.

All the H atoms were located from the difference Fourier map and were refined isotropically.

Data collection: *SMART* (Siemens, 1996); cell refinement: *SAINT* (Siemens, 1996); data reduction: *SAINT*; program(s) used to solve structure: *SHELXTL* (Sheldrick, 1997); program(s) used to refine

structure: *SHELXTL*; software used to prepare material for publication: *SHELXTL* and *PLATON* (Spek, 1990).

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