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Key indicators

Single-crystal X-ray study T = 295 KMean $\sigma(\text{C}-\text{C}) = 0.005 \text{ Å}$ R factor = 0.039 wR factor = 0.117 Data-to-parameter ratio = 20.4

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

[*N*,*N*'-Dimethyl-*N*"-(2-pyridylmethylene)ethane-1,2-diamine]dithiocyanatozinc(II)

The title compound, $[Zn(NCS)_2(C_{10}H_{15}N_3)]$, is a mononuclear zinc(II) complex. The molecule is isostructural with the analogous copper(II) compound [Yue, Xu, Shi & Feng (2005). *Acta Cryst.* E**61**, m693–m694]. The Zn atom exhibits a square pyramidal geometry.

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Comment

Zinc(II) complexes play an important role in numerous biological systems, where they usually function as the active site of hydrolytic enzymes (Casella & Gullotti, 1981; Leussing & Leach, 1971). As a part of our investigations of the structures of zinc derivatives, we have prepared the title compound, (I), a new mononuclear Zn(II) complex (Fig. 1), which is isostructural with the analogous copper(II) complex, [N,N-dimethyl-N'-(2-pyridylmethylene)ethane-1,2-diamino]dithiocyanatocopper(II), (Yue*et al.*, 2005). All the geometric parameters (Table 1) are within normal ranges (Erxleben, 2001). The Zn atom is five-coordinated by the three N atoms of the Schiff base and the terminal N atoms from two thiocyanate anions, giving a severely distorted square pyramidal geometry.



Experimental

N,N'-Dimethylethane-1,2-diamine (0.5 mmol, 44.1 mg) and pyridylaldehyde (0.5 mmol, 53.7 mg) were dissolved in EtOH (35 ml). The mixture was refluxed for 30 min, and aqueous solutions (5 ml) of ammonium thiocyanate (1.0 mmol, 76.2 mg) and Zn(ClO₄)₂·6H₂O (1.0 mmol, 372.4 mg) were added. The mixture was stirred under reflux for another 30 min to give a clear yellow solution. After allowing the solution to stand in air for 15 d, yellow crystals were formed.

Crystal data

$[Zn(NCS)_2(C_{10}H_{15}N_3)]$	$D_x = 1.461 \text{ Mg m}^{-3}$		
$M_r = 358.78$	Mo $K\alpha$ radiation		
Monoclinic, $P2_1/n$	Cell parameters from 5567		
a = 10.221 (1) Å	reflections		
b = 15.054 (2) Å	$\theta = 2.4-25.1^{\circ}$		
c = 10.600 (1) Å	$\mu = 1.76 \text{ mm}^{-1}$		
$\beta = 91.190 \ (1)^{\circ}$	T = 295 (2) K		
V = 1630.6 (3) Å ³ Block, yellow			
Z = 4	$0.22 \times 0.13 \times 0.11 \text{ mm}$		

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metal-organic papers

Data collection

Bruker APEX SMART CCD areadetector diffractometer3732 independ
2795 reflection ω scans2795 reflection ω scans $R_{int} = 0.028$ Absorption correction: multi-scan
(SADABS; Sheldrick, 1996) $\theta_{max} = 27.5^{\circ}$ $T_{min} = 0.698, T_{max} = 0.830$ $k = -13 \rightarrow 13$ 18260 measured reflections $l = -13 \rightarrow 13$ RefinementRefinement

Refinement of F $R[F^2 > 2\sigma(F^2)] = 0.039$ $wR(F^2) = 0.117$ S = 1.03 3732 reflections 183 parameters H-atom parameters constrained 3732 independent reflections 2795 reflections with $I > 2\sigma(I)$ $R_{int} = 0.028$ $\theta_{max} = 27.5^{\circ}$ $h = -13 \rightarrow 13$ $k = -19 \rightarrow 19$

$$\begin{split} &w = 1/[\sigma^2(F_{\rm o}^2) + (0.0616P)^2 \\ &+ 0.5292P] \\ &where \ P = (F_{\rm o}^2 + 2F_{\rm c}^2)/3 \\ (\Delta/\sigma)_{\rm max} < 0.001 \\ \Delta\rho_{\rm max} = 1.07 \ {\rm e} \ {\rm \AA}^{-3} \\ \Delta\rho_{\rm min} = -0.17 \ {\rm e} \ {\rm \AA}^{-3} \end{split}$$

Table 1

Selected geometric parameters (Å, °).

Zn1-N5	1.962 (3)	Zn1-N1	2.207 (3)
Zn1-N4	1.964 (3)	Zn1-N3	2.224 (3)
Zn1-N2	2.061 (2)		
N5-Zn1-N4	111.14 (14)	N2-Zn1-N1	75.35 (10)
N5-Zn1-N2	135.41 (13)	N5-Zn1-N3	96.65 (11)
N4-Zn1-N2	113.31 (12)	N4-Zn1-N3	101.36 (11)
N5-Zn1-N1	96.39 (11)	N2-Zn1-N3	78.16 (10)
N4-Zn1-N1	96.59 (11)	N1-Zn1-N3	152.25 (9)

All H atoms were placed in idealized positions and constrained to ride on their parent atoms, with C-H distances in the range 0.93–0.97 Å, and with $U_{iso}(H)$ set to 1.2 or 1.5 times $U_{eq}(C)$. The highest electron density peak, 1.07 e Å⁻³, is located 1.14 Å from atom N1, but no physical significance can be attached to this observation.

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINT* (Bruker, 1998); 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*.



Figure 1

The structure of (I). Displacement ellipsoids are drawn at the 30% probability level.

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