

*N,N'*-Bis[2-(2-pyridyl)ethyl]dithiooxamideJian-Zhong Cui,<sup>a\*</sup> Dan Zhang,<sup>a</sup>  
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## Key indicators

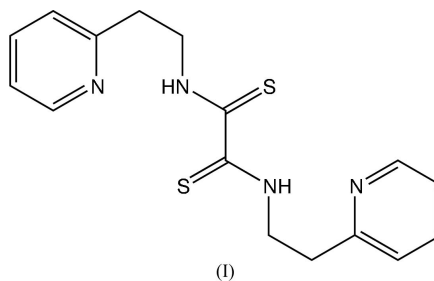
Single-crystal X-ray study  
*T* = 293 K  
Mean  $\sigma(\text{C}-\text{C})$  = 0.002 Å  
*R* factor = 0.033  
*wR* factor = 0.097  
Data-to-parameter ratio = 19.1For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.

The title compound,  $\text{C}_{16}\text{H}_{18}\text{N}_4\text{S}_2$ , has a crystallographic inversion centre at the mid-point of the central C—C bond. The plane of the dithiooxamide fragment is approximately perpendicular to the plane of the dimethylene bridge [dihedral angle =  $84.8(1)^\circ$ ], the plane of which is in turn normal to the pyridine ring plane [dihedral angle =  $89.80(8)^\circ$ ]. An intermolecular N—H $\cdots$ N hydrogen bond involving the amide NH group and the pyridine N atom [ $\text{N}\cdots\text{N}$  =  $3.0503(18)$  Å and  $\text{N}-\text{H}\cdots\text{N}$  =  $145.8(14)^\circ$ ] links the molecules into chains extending along the *c* axis.

## Comment

The title compound is a representative of the relatively scarcely studied *N*-substituted thiooxamides, which may be used as versatile nitrogen- and sulfur-containing ligands for multi-nuclear metal complexes with special magnetic properties (Hurd *et al.*, 1960). Only a few such molecules have been reported (Cui *et al.*, 2004), because of the difficulties in their preparation and purification.

The title compound, (I), was synthesized according to the method of Cui *et al.* (2004); its molecular structure is shown in Fig. 1. The molecule of (I) has a crystallographic inversion centre at the mid-point of the C1—C1A bond. The dithiooxamide plane, defined by atoms N1, C1 and S1 and their symmetry-related counterparts, is approximately orthogonal to the mean plane of the dimethylene bridge, N1/C2—C4 [the dihedral angle is  $84.8(1)^\circ$ ], which in turn is almost orthogonal to the pyridine plane, N2/C4—C8 [dihedral angle  $89.80(8)^\circ$ ]. The pyridine plane forms a small dihedral angle of  $6.2(2)^\circ$  with the central dithiooxamide plane.



An intermolecular N1—H1 $\cdots$ N2<sup>ii</sup> bond [symmetry code: (ii)  $-x + 1, y - \frac{1}{2}, -z + \frac{3}{2}$ ; Table 2] links the molecules of (I) into chains extending along the *c* axis. The packing of the molecules in the crystal structure of (I) is shown in Fig. 2.

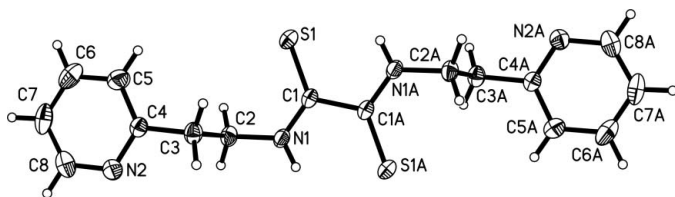
## Experimental

The title compound was prepared as an orange powder, with a melting point of 431 K, according to the procedure reported by Cui *et*

Received 30 March 2005

Accepted 22 May 2005

Online 28 May 2005



**Figure 1**  
View of the molecule, showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 30% probability level. H atoms are shown as circles of arbitrary size. [Symmetry code: (A)  $1 - x, -y, 1 - z$ ].

*al.* (2004) (yield 80%). Single crystals suitable for X-ray diffraction study were obtained by slow evaporation of an ethanol solution.

**Crystal data**

$C_{16}H_{18}N_4S_2$   $D_x = 1.317 \text{ Mg m}^{-3}$   
 $M_r = 330.46$  Mo  $K\alpha$  radiation  
 Monoclinic,  $P2_1/c$  Cell parameters from 1570 reflections  
 $a = 10.4499 (18) \text{ \AA}$   
 $b = 6.0036 (10) \text{ \AA}$   
 $c = 13.688 (2) \text{ \AA}$   
 $\beta = 103.995 (2)^\circ$   
 $V = 833.2 (2) \text{ \AA}^3$   
 $Z = 2$   $\theta = 3.2\text{--}27.1^\circ$   
 $\mu = 0.32 \text{ mm}^{-1}$   
 $T = 293 (2) \text{ K}$   
 Prism, orange  
 $0.28 \times 0.24 \times 0.20 \text{ mm}$

**Data collection**

Bruker SMART CCD area-detector diffractometer 1982 independent reflections  
 $\varphi$  and  $\omega$  scans 1491 reflections with  $I > 2\sigma(I)$   
 Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996)  $R_{int} = 0.018$   
 $T_{min} = 0.902, T_{max} = 0.938$   $\theta_{max} = 27.9^\circ$   
 5351 measured reflections  $h = -13 \rightarrow 10$   
 $l = -15 \rightarrow 18$

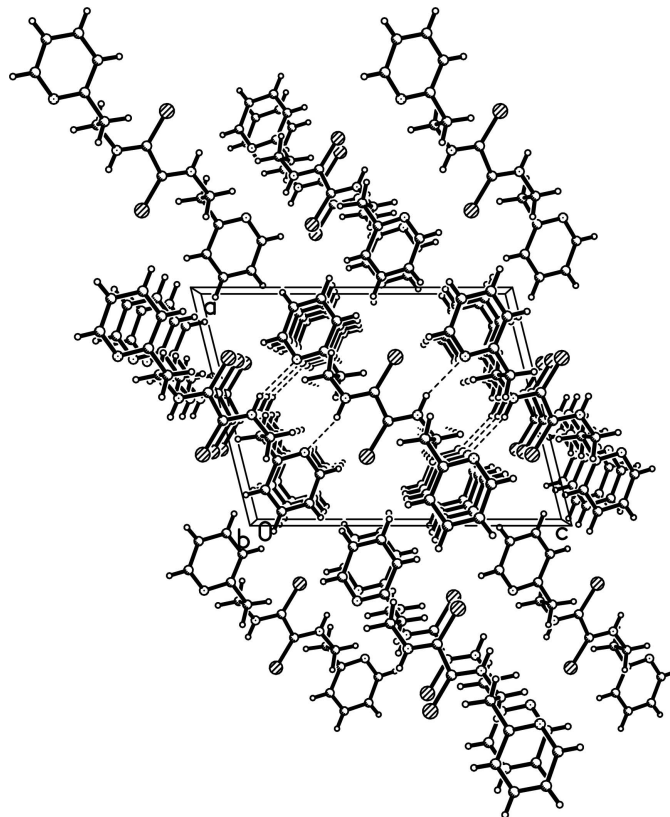
**Refinement**

Refinement on  $F^2$  H atoms treated by a mixture of independent and constrained refinement  
 $R[F^2 > 2\sigma(F^2)] = 0.033$   
 $wR(F^2) = 0.097$   
 $S = 1.11$   
 1982 reflections  
 104 parameters  
 $w = 1/[\sigma^2(F_o^2) + (0.0729P)^2]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{max} = 0.001$   
 $\Delta\rho_{max} = 0.23 \text{ e \AA}^{-3}$   
 $\Delta\rho_{min} = -0.16 \text{ e \AA}^{-3}$

**Table 1**  
Selected geometric parameters ( $\text{\AA}, ^\circ$ ).

S1—C1	1.6591 (13)	C2—C3	1.520 (2)
N1—C1	1.3112 (18)	C3—C4	1.498 (2)
N1—C2	1.4511 (17)	C4—C5	1.377 (2)
N2—C4	1.3256 (19)	C5—C6	1.367 (3)
N2—C8	1.338 (2)	C6—C7	1.362 (3)
C1—C1 <sup>1</sup>	1.530 (2)	C7—C8	1.366 (3)
C1—N1—C2	124.12 (12)	N2—C4—C5	121.91 (14)
C4—N2—C8	117.21 (14)	N2—C4—C3	117.04 (13)
N1—C1—C1 <sup>1</sup>	114.32 (14)	C5—C4—C3	121.01 (13)
N1—C1—S1	124.50 (10)	C6—C5—C4	119.75 (17)
C1 <sup>1</sup> —C1—S1	121.18 (13)	C7—C6—C5	119.06 (17)
N1—C2—C3	112.12 (12)	C6—C7—C8	117.93 (17)
C4—C3—C2	110.79 (12)	N2—C8—C7	124.15 (18)
C2—N1—C1—C1 <sup>1</sup>	-179.72 (13)	C2—C3—C4—N2	89.18 (16)
C2—N1—C1—S1	0.23 (19)	C2—C3—C4—C5	-88.52 (18)
C1—N1—C2—C3	-84.01 (17)	C3—C4—C5—C6	177.57 (16)
N1—C2—C3—C4	175.80 (12)	C4—C5—C6—C7	0.3 (3)
C8—N2—C4—C5	-0.3 (2)	C6—C7—C8—N2	-0.1 (3)
C8—N2—C4—C3	-177.96 (13)		

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



**Figure 2**  
Packing of the molecules in the crystal structure of (I), viewed down the *b* axis. Hydrogen bonds are shown as dashed lines.

**Table 2**  
Hydrogen-bond geometry ( $\text{\AA}, ^\circ$ ).

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
N1—H1...N2 <sup>ii</sup>	0.85 (1)	2.31 (1)	3.0503 (18)	146 (1)

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

The H atom of the NH group was located in a difference Fourier map and refined isotropically, subject to the restraint  $N-H = 0.85 (1) \text{ \AA}$ . All C-bound H atoms were positioned geometrically and included in the refinement in the riding-model approximation ( $C-H = 0.93\text{--}0.97 \text{ \AA}$ ).  $U_{iso}(H)$  values were set to  $1.2U_{eq}(\text{carrier atom})$ .

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

This work was supported by the Scientific Research Foundation for Returned Overseas Chinese Scholars, the Ministry of Education, People's Republic of China.

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