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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.004 Å R factor = 0.041 wR factor = 0.093 Data-to-parameter ratio = 12.5

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

A three-dimensional hydrogen-bonded network of diaquabis(2,5-di-3-pyridyl-1,3,4-oxadiazole)dithiocyanatoiron(II)

The molecular structure of the neutral mononuclear complex $[Fe(NCS)_2(3-bpo)_2(H_2O)_2]$, in which 3-bpo denotes 2,5-di-3pyridyl-1,3,4-oxadiazole $(C_{12}H_8N_4O)$, is centrosymmetric, with Fe^{II} located on an inversion center. This complex is octahedral, and the Fe^{II} atom is *trans* coordinated by two monodentate NSC⁻ ligands, two monodentate 3-bpo ligands and two water molecules. Each complex is linked to four others, forming a (011) sheet, by O-H…N hydrogen bonding. Weak C-H…S interactions link each (011) sheet to the two adjacent sheets, hence forming a three-dimensional array. There is also some slipped π stacking within each sheet.

Comment

This work is part of our continuing interest in metalcontaining assemblies based on dipyridyl ligands containing the oxadiazole spacer, such as 2,5-di-4-pyridyl-1,3,4-oxadiazole (4-bpo) and its 3,3'-*N*-donor analog (3-bpo), *via* coordination, hydrogen-bonded and/or other intermolecular cooperative interactions (Du, Zhao & Guo, 2005, and references therein; Du & Zhao, 2004). Very recently, we have described an Mn^{II} -3-bpo–NCS mononuclear complex, [$Mn(NCS)_2(3-bpo)_2(H_2O)_2$], with a three-dimensional hydrogen-bonded network (Du, Jiang & Zhao, 2005), and we report here the isostructural Fe^{II} compound, namely the title complex, (I).



In the molecular structure (Fig. 1) of the neutral mononuclear complex (I), each Fe^{II} atom, located on an inversion center, is six-coordinated by two monodentate NSC⁻ anions and two water molecules, forming the equatorial plane, and

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Figure 1

The title complex, with atom labeling of the asymmetric unit and the Fe^{II} coordination sphere, shown with 30% probability displacement ellipsoids. [Symmetry code: (A) 1 - x, 1 - y, 1 - z.]

two pyridyl N atoms of a pair of monodentate 3-bpo ligands in the axial positions, displaying an FeN₄O₂ octahedral geometry. The metal-ligand distances within the octahedral framework are close to values reported for related $[Fe(NCS)_2(L)_2(H_2O)_2]$ complexes (Long et al., 1983; Cingi et al., 1986; Noro et al., 1999). The 3-bpo ligand is not planar; the pyridyl rings make dihedral angles of 6.8 (2) and 10.0 (2) $^{\circ}$ with the central oxadiazole system, and the dihedral angle between them is 5.0 (2)°.

The 3-bpo molecule takes the unusual trans configuration with respect to the two terminal pyridyl groups (Du & Zhao, 2003), in order to favor the hydrogen bond between the uncoordinated 3-pyridyl N atom and the aqua ligand. Each water molecule also generates a hydrogen bond with one N atom of the oxadiazole group. Thus, each complex is linked to four others, forming a (011) sheet, by $O-H \cdots N$ hydrogen bonds. Weak $C-H \cdots S$ interactions link each (011) sheet to the two adjacent sheets, hence forming a three-dimensional array (Table 1). There is also some slipped π stacking within each sheet. The centroid-to-centroid and centroid-to-plane distances between two neighboring almost antiparallel pyridyl rings are 3.87 and 3.50/3.42 Å, respectively.

Experimental

Fe(ClO₄)₂·6H₂O (0.1 mmol), 3-bpo (0.1 mmol) and NH₄SCN (0.1 mmol) were mixed in a CH₃CN/H₂O (20 ml, 1:1) solution with refluxing for ca 20 min. The resulting solution was filtered and left to stand at room temperature. Red prismatic crystals suitable for X-ray analysis were obtained in 50% yield by slow evaporation of the solvent over a period of 1 week. Analysis calculated for C₂₆H₂₀FeN₁₀O₄S₂: C 47.57, H 3.07, N 21.34%; found: C 47.55, H 2.98, N 21.00%. IR (KBr pellet, cm⁻¹): 3235 (*b*), 3070 (*m*), 2079 (*vs*), 1663 (m), 1606 (vs), 1587 (s), 1551 (m), 1485 (s), 1464 (s), 1436 (s), 1417 (s),1337 (m), 1276 (m), 1196 (m), 1128 (m), 1116 (m), 1087 (s), 1042 (s), 1028 (s), 997 (w), 965 (m), 823 (s), 733 (s), 699 (vs), 636 (m).

Crystal data

$[Fe(C_{12}H_8N_4O)_2(NCS)_2(H_2O)_2]$	Z = 1
$M_r = 656.49$	$D_x = 1.522 \text{ Mg m}^{-3}$
Triclinic, P1	Mo $K\alpha$ radiation
a = 8.159 (7) Å	Cell parameters from 697
b = 8.712 (7) Å	reflections
c = 10.531 (9) Å	$\theta = 2.4 - 26.0^{\circ}$
$\alpha = 82.385 \ (13)^{\circ}$	$\mu = 0.73 \text{ mm}^{-1}$
$\beta = 77.494 \ (13)^{\circ}$	T = 293 (2) K
$\gamma = 80.064 \ (14)^{\circ}$	Prism, red
$V = 716.3 (10) \text{ Å}^3$	$0.20 \times 0.18 \times 0.14 \text{ mm}$



Figure 2

The two-dimensional supramolecular structure of the title complex, formed via O-H···N hydrogen-bonds between water and 3-bpo (indicated by purple dashed lines).

Data collection

Bruker APEX II CCD area-	2517 independent reflections
detector diffractometer	1851 reflections with $I > 2\sigma(I)$
φ and ω scans	$R_{\rm int} = 0.023$
Absorption correction: multi-scan	$\theta_{\rm max} = 25.0^{\circ}$
(SADABS; Sheldrick, 1996)	$h = -9 \rightarrow 6$
$T_{\min} = 0.810, \ T_{\max} = 0.903$	$k = -10 \rightarrow 9$
3735 measured reflections	$l = -12 \rightarrow 8$

Refinement

Refinement on F^2
$R[F^2 > 2\sigma(F^2)] = 0.041$
$\nu R(F^2) = 0.093$
1 = 1.02

2517 reflections

202 parameters

H atoms treated by a mixture of independent and constrained refinement

$R_{\rm int} = 0.023$	
$\theta_{\rm max} = 25.0^{\circ}$	
$h = -9 \rightarrow 6$	
$k = -10 \rightarrow 9$	
$l = -12 \rightarrow 8$	

$w = 1/[\sigma^2(F_o^2) + (0.0371P)^2]$
+ 0.299P]
where $P = (F_0^2 + 2F_c^2)/3$
$(\Delta/\sigma)_{\rm max} < 0.001$
$\Delta \rho_{\rm max} = 0.33 \text{ e } \text{\AA}^{-3}$
$\Delta \rho_{\rm min} = -0.31 \text{ e } \text{\AA}^{-3}$

Table 1		
Hydrogen-bond geometry	(Å,	°).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$\begin{array}{c} O2 - H2A \cdots N3^{i} \\ O2 - H2B \cdots N4^{ii} \end{array}$	0.84(1)	1.98 (1)	2.807 (4)	167 (3)
	0.84(1)	1.95 (1)	2.775 (3)	168 (3)
$\begin{array}{l} \text{C10-H10} \cdots \text{S1}^{\text{iii}} \\ \text{C9-H9} \cdots \text{S1}^{\text{iv}} \end{array}$	0.93	2.83	3.575 (4)	138
	0.93	2.85	3.625 (4)	141

Symmetry codes: (i) -x + 1, -y + 1, -z; (ii) x, y - 1, z + 1; (iii) x + 1, y + 1, z - 1; (iv) x + 1, -v + 1, -z + 1.

H atoms were located in difference maps, but those attached to C atoms were introduced in calculated positions and treated as riding, with C-H = 0.93Å. H atoms of water molecules were refined using restraints $[O-H = 0.85 (1) \text{ Å and } H \cdots H = 1.39 (2) \text{ Å}]$. All H atoms were allocated displacement parameters related to those of their parent atoms $[U_{iso}(H) = 1.2U_{eq}(C,O)].$

Data collection: APEX2 (Bruker, 2003); cell refinement: APEX2; data reduction: SAINT (Bruker, 2001); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Bruker, 2001) and DIAMOND (Brandenburg & Berndt, 1999); software used to prepare material for publication: SHELXTL.

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