Received 23 November 2004

Accepted 6 December 2004

Online 11 December 2004

Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

Gregory J. Halder and Cameron J. Kepert*

School of Chemistry, University of Sydney, NSW 2006, Australia

Correspondence e-mail: c.kepert@chem.usyd.edu.au

Key indicators

Single-crystal X-ray study T = 150 K Mean σ (C–C) = 0.002 Å R factor = 0.023 wR factor = 0.064 Data-to-parameter ratio = 14.0

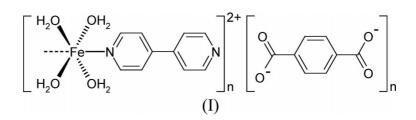
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

A three-dimensional hydrogen-bonded coordination framework: *catena*-poly[[[tetraaquairon(II)]- μ -4,4'-bipyridine] benzene-1,4-dicarboxylate]

The title compound, {[Fe(C₁₀H₈N₂)(H₂O)₄](C₈H₄O₄)]_n or [Fe(bpy)(H₂O)₄](bdc) (bpy is 4,4'-bipyridine and bdc is benzene-1,4-dicarboxylate), contains one-dimensional chains of Fe^{II} centres bridged by bpy ligands along the *b* axis, with the equatorial coordination of four water molecules completing the octahedral coordination. Non-coordinated bdc anions balance the charge of the Fe^{II} centres and bridge adjacent one-dimensional chains through an extensive hydrogen-bonding network. The compound crystallizes in the centrosymmetric space group P2/c, which defines twofold axes along both the Fe–bpy chains and the bdc anions.

Comment

The linking of metal ions with bridging organic ligands offers a versatile means of constructing materials with interesting structure and function (Hoskins & Robson, 1990). The linear bridging ligands 4,4'-bipyridine (bpy) and benzene-1,4-dicarboxylate (bdc) have been extensively utilized in the formation of such materials (*e.g.* Kepert & Rosseinsky, 1999; Li *et al.*, 1999). The present study of the title compound, (I), was carried out with the aim of engineering a three-dimensional coordination framework of (4,4) Fe–bpy sheets bridged by bdc pillars. However, the incorporation of coordinated water molecules resulted in a hydrogen-bonded network.



Compound (I) crystallizes in the centrosymmetric monoclinic space group P2/c. The primary structure consists of onedimensional chains of Fe^{II} centres bridged by bpy ligands along the *b* axis. The equatorial coordination of four water molecules completes the mildly distorted octahedral Fe^{II} coordination. The Fe–O and Fe–N bond lengths are consistent with high-spin Fe^{II} (Table 1).

Non-coordinated bdc ligands balance the charge of the system and bridge adjacent one-dimensional chains through an extensive hydrogen-bonding network (Fig. 2). All water H atoms form hydrogen bonds with either of the two crystal-lographically distinct carboxylate O atoms (Table 2). These interactions are likely to contribute to the highly ordered nature of the structure.

© 2005 International Union of Crystallography Printed in Great Britain – all rights reserved

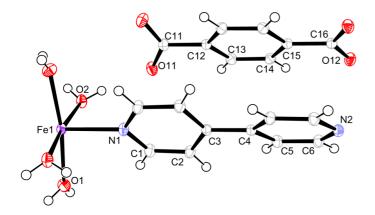


Figure 1

A view of the structure of (I), showing 50% probability displacement ellipsoids and the atom-numbering scheme. Unlabelled atoms are related to labelled atoms by the symmetry operation $(1 - x, y, \frac{3}{2} - z)$.

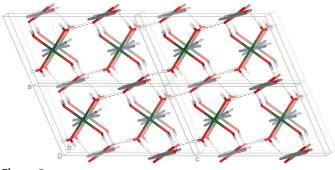


Figure 2

The crystal packing of (I) viewed down the b axis, showing the extensive hydrogen-bonding network.

Experimental

 $[Fe(bpy)(H_2O)_4](bdc)$, (I), was synthesized by the slow diffusion of aqueous solutions of Fe(ClO₄)₂ (31 mg, 0.12 mmol) and a mixture of bpy (37 mg, 0.24 mmol) and Na₂bdc (25 mg, 0.12 mmol) in an Hshaped tube. Orange block-shaped single crystals of (I) grew after approximately one month.

Crystal data

$[Fe(C_{10}H_8N_2)(H_2O)_4](C_8H_4O_4)$ $M_r = 448.21$ Monoclinic, $P2/c$ a = 6.9541 (12) Å b = 11.465 (2) Å c = 12.059 (2) Å $\beta = 113.547$ (3)° V = 881.4 (3) Å ³ Z = 2	$D_x = 1.689 \text{ Mg m}^{-3}$ Mo K\alpha radiation Cell parameters from 889 reflections $\theta = 3.2-27.9^{\circ}$ $\mu = 0.91 \text{ mm}^{-1}$ T = 150 (2) K Block, orange $0.50 \times 0.37 \times 0.22 \text{ mm}$
Data collection	
Bruker SMART 1000 CCD area- detector diffractometer ω scans Absorption correction: multi-scan, (<i>SADABS</i> ; Sheldrick, 1996) $T_{\min} = 0.733, T_{\max} = 0.819$ 8043 measured reflections	2075 independent reflections 2005 reflections with $I > 2\sigma(I)$ $R_{int} = 0.017$ $\theta_{max} = 28.0^{\circ}$ $h = -9 \rightarrow 8$ $k = -15 \rightarrow 15$ $l = -15 \rightarrow 15$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0212P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.023$	+ 0.3552P]
$wR(F^2) = 0.065$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.13	$(\Delta/\sigma)_{\rm max} < 0.001$
2075 reflections	$\Delta \rho_{\rm max} = 0.26 \ {\rm e} \ {\rm \AA}^{-3}$
148 parameters	$\Delta \rho_{\rm min} = -0.44 \ {\rm e} \ {\rm \AA}^{-3}$
H atoms treated by a mixture of	
independent and constrained	
refinement	

Table 1

Selected geometric parameters (Å, °).

Fe1-O1	2.1311 (10)	Fe1-N1	2.1495 (15)
Fe1-O2	2.1375 (10)	Fe1-N2 ⁱ	2.2081 (14)
O1-Fe1-O1 ⁱⁱ	171.39 (5)	O2-Fe1-N1	86.32 (2)
O1-Fe1-O2 ⁱⁱ	86.77 (4)	O1-Fe1-N2 ⁱ	85.69 (3)
O1-Fe1-O2	93.79 (4)	O2-Fe1-N2 ⁱ	93.68 (2)
O2 ⁱⁱ -Fe1-O2	172.65 (5)	N1-Fe1-N2 ⁱ	180.0
O1-Fe1-N1	94.31 (3)		

Symmetry codes: (i) x, y + 1, z; (ii) $-x + 1, y, -z + \frac{3}{2}$.

Table 2 Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
$01 - H1A \cdots O12^{iii}$ $01 - H1B \cdots O11^{iv}$	0.797 (13)	2.074 (14)	2.8578 (13)	167.7 (17)
	0.813 (13)	1.910 (14)	2.7204 (14)	174.3 (17)
$\begin{array}{c} O2 - H2A \cdots O11^{v} \\ O2 - H2B \cdots O12 \end{array}$	0.796 (13)	1.965 (13)	2.7570 (13)	174.0 (17)
	0.816 (13)	1.907 (13)	2.7220 (13)	177.2 (18)

Symmetry codes: (iii) x + 1, y, z; (iv) -x + 1, -y + 2, -z + 2; (v) x, y - 1, z.

All H atoms were modelled with fixed isotropic parameters of $1.2U_{eq}$ (aromatic) or $1.5U_{eq}$ (water) of their parent atom. Aromatic H-atom sites were constrained to geometrically idealized positions, with C-H distances of 0.95 Å. Water H atoms were located from a difference map and the O-H bond lengths were restrained to be approximately equal.

Data collection: SMART (Bruker, 1997); cell refinement: SAINT-Plus (Bruker, 1997); data reduction: SAINT-Plus; program(s) used to solve structure: SHELXTL (Bruker, 2000); program(s) used to refine structure: SHELXTL; molecular graphics: ORTEP-3 (Farrugia, 1997) and WebLab ViewerPro3.7 (Accelrys, 2001); software used to prepare material for publication: XCIF in SHELXTL.

The authors thank the Australian Research Council for financial support.

References

Accelrys (2001). WebLab ViewerPro3.7. Accelrys Inc., San Diego, California, USA.

Bruker (1997). SMART (Version 5.504) and SAINT-Plus (Version 6.45). Bruker AXS Inc., Madison, Wisconsin, USA.

Bruker (2000). SHELXTL. Version 6.14. Bruker AXS Inc., Madison, Wisconsin, USA.

Farrugia, L. J. (1997). J. Appl. Cryst. 30, 565.

Hoskins, B. F. & Robson, R. (1990). J. Am. Chem. Soc. 112, 1546-1554.

Kepert, C. J. & Rosseinsky, M. J. (1999). Chem. Commun. pp. 375-376.

Li, H., Eddaoudi, M., O'Keeffe, M. & Yaghi, O. (1999). Nature (London), 402, 276-279.

Sheldrick, G. M. (1996). SADABS. University of Göttingen, Germany.