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Key indicators

Single-crystal X-ray study
 T = 293 K
 Mean $\sigma(\text{C}-\text{C}) = 0.007 \text{ \AA}$
 R factor = 0.052
 wR factor = 0.139
 Data-to-parameter ratio = 11.7

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

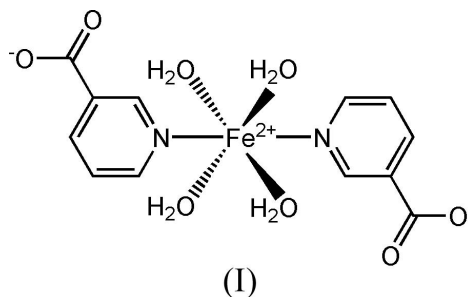
Tetraaquabis(pyridine-3-carboxylato- κN)iron(II)

Pyridine-3-carboxylic acid (nicH) reacts with iron(II) chloride tetrahydrate in a methanol–water mixture to form the title compound, $[\text{Fe}(\text{C}_6\text{H}_4\text{NO}_2)_2(\text{H}_2\text{O})_4]$. The Fe^{II} atom is in a slightly distorted octahedral coordination. The crystal structure is stabilized by $\text{O}-\text{H}\cdots\text{O}$ hydrogen bonds involving all the water H atoms and the carboxylate groups.

Received 11 July 2005
 Accepted 9 August 2005
 Online 17 August 2005

Comment

Pyridinecarboxylates and their derivatives are useful ligands in the construction of one- to three-dimensional functional metal–organic frameworks (Evans & Lin, 2002; Lu, 2003; Tong *et al.*, 2003, 2004, 2005; Wang *et al.*, 2003). The title complex, (I), was obtained in an attempt to prepare a multidimensional coordination network.



In the structure of (I), the Fe^{II} atom, which occupies a special position of $2/m$ symmetry, is coordinated by four aqua ligands and two pyridyl N atoms from two pyridine-3-carboxylate (nic) ligands (Fig. 1). The $\text{Fe1}-\text{O}$ [2.210 (3) \AA]

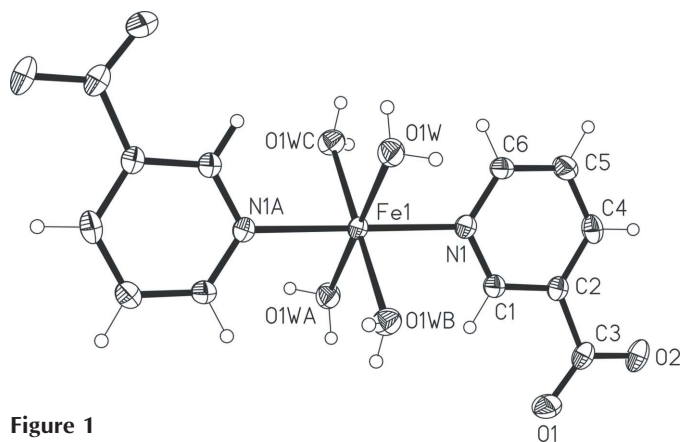


Figure 1 ORTEP plot (Johnson, 1976) of the molecular structure of (I), showing the atom-labelling scheme. Displacement ellipsoids are plotted at the 40% probability level and H atoms are shown as small spheres of arbitrary radii. [Symmetry codes: (A) $x, -y, z$; (B) $-x, y, -z$; (C) $-x, -y, -z$.]

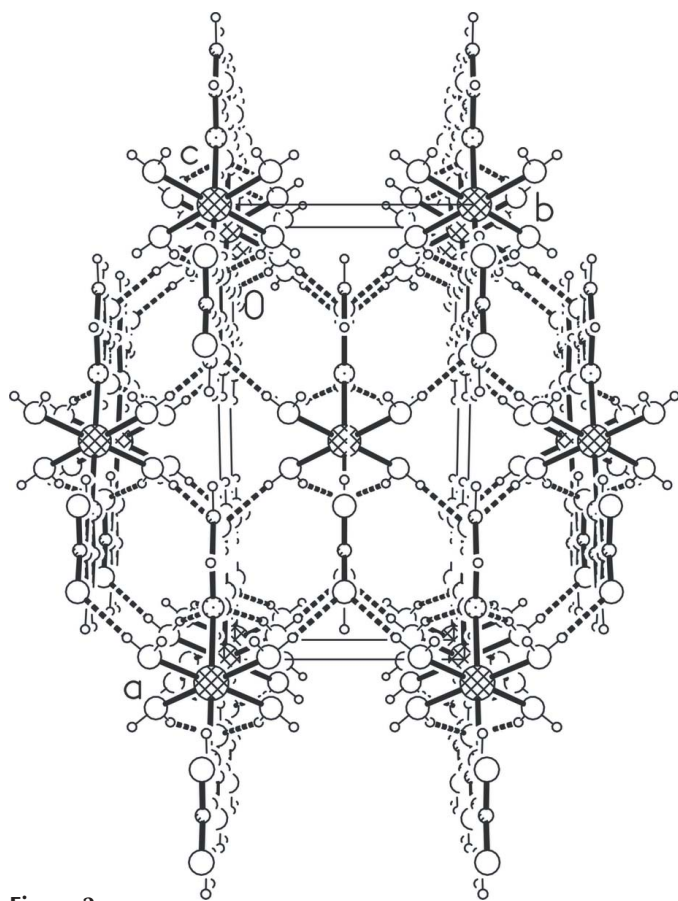


Figure 2
Perspective view of the molecular packing of (I).

and Fe1–N [2.135 (2) Å] bond lengths are similar to those found in tetraaquabis(pyridine-4-carboxylato- κ N)iron(II) [Fe–N = 2.202 (2) Å and Fe–O = 2.125 (2) and 2.136 (2) Å; Liu *et al.*, 1999]. The Fe^{II} atom, the pyridyl rings and the carboxylate groups are coplanar. The structure of (I) is isostructural with that of *trans*-tetraaquabis(pyridine-3-carboxylato- κ N)chromium(II) (Cotton *et al.*, 1991), *trans*-tetraaquabis(pyridine-3-carboxylato- κ N)cobalt(II) (Anagnostopoulos *et al.*, 1969), *trans*-tetraaquabis(pyridine-3-carboxylato- κ N)nickel(II) (Batten & Harris, 2001), *trans*-tetraaquabis(pyridine-3-carboxylato- κ N)zinc(II) (Cingi *et al.*, 1971), and *trans*-tetraaquabis(pyridine-3-carboxylato- κ N)cadmium(II) (Zhou *et al.*, 2003). The extensive intermolecular O–H...O hydrogen bonds [O...O = 2.696 (4)–2.713 (3) Å] between the carboxylate groups and the aqua ligands play an important role in consolidating the crystal architecture (Fig. 2).

Experimental

To a methanol–water solution (5:1 *v/v*), 10 ml of iron(II) chloride tetrahydrate (0.100 g, 0.5 mmol), a solution of pyridine-3-carboxylic acid (0.123 g, 1.0 mmol) in MeOH (5 ml) was added slowly, with stirring, over a period of 10 min at ambient temperature. Red block-shaped crystals were deposited within 2 d in *ca* 45% yield.

Crystal data

[Fe(C₆H₄NO₂)₂(H₂O)₄]
M_r = 372.12
 Monoclinic, *C*2/*m*
a = 14.235 (4) Å
b = 6.901 (2) Å
c = 8.538 (3) Å
 β = 118.12 (1)°
V = 739.7 (4) Å³
Z = 2

D_x = 1.671 Mg m⁻³
 Mo *K* α radiation
 Cell parameters from 680 reflections
 θ = 3.3–27.5°
 μ = 1.06 mm⁻¹
T = 293 (2) K
 Block, red
 0.32 × 0.26 × 0.18 mm

Data collection

Bruker SMART CCD area-detector diffractometer
 φ and ω scans
 Absorption correction: multi-scan (SADABS; Blessing, 1995)
T_{min} = 0.704, *T_{max}* = 0.821
 2628 measured reflections

886 independent reflections
 770 reflections with *I* > 2 σ (*I*)
R_{int} = 0.000
 θ_{\max} = 27.5°
h = -18 → 13
k = -7 → 8
l = -10 → 11

Refinement

Refinement on *F*²
R[*F*² > 2 σ (*F*²)] = 0.052
wR(*F*²) = 0.139
S = 1.03
 886 reflections
 76 parameters

H atoms treated by a mixture of independent and constrained refinement
 $w = 1/[\sigma^2(F_o^2) + (0.097P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.72 \text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.73 \text{ e \AA}^{-3}$

Table 1

Selected geometric parameters (Å, °).

Fe1–O1W	2.135 (2)	Fe1–N1	2.210 (3)
O1W ⁱ –Fe1–O1W	90.59 (14)	O1W–Fe1–N1	90.21 (9)
O1W–Fe1–O1W ⁱⁱ	89.41 (14)	C5–C6–H6	118.3
O1W–Fe1–N1 ⁱⁱⁱ	89.79 (9)		

Symmetry codes: (i) *x*, -*y*, *z*; (ii) -*x*, *y*, -*z*; (iii) -*x*, -*y*, -*z*.

Table 2

Hydrogen-bond geometry (Å, °).

<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>
O1W–H1WA...O1 ^{iv}	0.85 (1)	1.88 (1)	2.696 (4)	161 (4)
O1W–H1WB...O2 ^v	0.85 (1)	1.87 (1)	2.713 (3)	174 (4)

Symmetry codes: (iv) *x*, *y*, *z* - 1; (v) -*x* + $\frac{1}{2}$, -*y* + $\frac{1}{2}$, -*z* + 1.

The pyridyl H atoms were included in geometrically calculated positions and constrained to ride, with C–H = 0.93 Å and *U*_{iso}(H) = 1.2*U*_{eq}(C). The aqua H atoms were located in difference Fourier maps and refined with isotropic displacement parameters subject to the restraint O–H = 0.85 (1) Å.

Data collection: SMART (Bruker, 1998); cell refinement: SMART; data reduction: SAINT (Bruker, 1999) and SHELXTL (Bruker, 1998); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL; software used to prepare material for publication: SHELXL97.

The authors thank the Foundation for the Items of Education Office of Guangdong Province (No, z03065) for generously supporting this work.

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