

Tris(1,10-phenanthroline- κ^2N,N')iron(III) trinitrate monohydrateMamiko Odoko* and
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Key indicators

Single-crystal X-ray study
 $T = 296$ K
Mean $\sigma(C-C) = 0.004$ Å
Disorder in solvent or counterion
 R factor = 0.043
 wR factor = 0.123
Data-to-parameter ratio = 15.1For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.

In the title compound, $[\text{Fe}(\text{C}_{12}\text{H}_8\text{N}_2)_3](\text{NO}_3)_3 \cdot \text{H}_2\text{O}$, the Fe atom lies on a twofold rotation axis and is coordinated by three planar 1,10-phenanthroline ligands in a slightly distorted octahedral environment. One nitrate anion and the disordered water molecule also lie on twofold rotation axes. The crystal structure is stabilized by partial stacking and hydrogen-bonding interactions.

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Comment

1,10-Phenanthroline (phen) molecules can stack with nucleobases to enhance complex formation with DNA, which is the principal target in the chemotherapy of tumors. Fe complexes containing phen have been studied for their ability to interact with DNA (Furtado *et al.*, 1997; Mudasir *et al.*, 2003). Few crystal structures of Fe^{3+} complexes of phen have been reported (Baker *et al.*, 1975), although there are many papers describing crystal structures containing $[\text{Fe}(\text{phen})_3]^{2+}$ cations (Broomhead *et al.*, 1982; Che *et al.*, 1997; Deng *et al.*, 2001; Hoshina *et al.*, 2000; Koh *et al.*, 1994; Li *et al.*, 1993; Reiff *et al.*, 1983; Ronco *et al.*, 1986; Yan *et al.*, 2000; Zalkin *et al.*, 1973). In the present study, we synthesized the title compound, $[\text{Fe}(\text{phen})_3]^{3+} \cdot 3\text{NO}_3^- \cdot \text{H}_2\text{O}$, and analyzed its crystal structure.

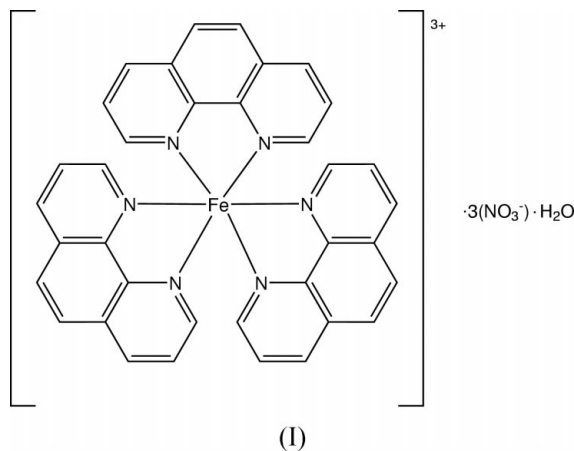


Fig. 1 shows a view of the $[\text{Fe}(\text{phen})_3]^{3+}$ cation. The Fe atom, which lies on a twofold rotation axis, is coordinated by six N atoms of three phen ligands to form a distorted octahedral environment. The Fe1–N1 bond length, 1.988 (1) Å, is slightly longer than the other Fe–N bonds; the average Fe–N bond length is 1.976 Å, in agreement with previously reported results for the $[\text{Fe}(\text{phen})_3]^{3+}$ cation (Baker *et al.*, 1975), ranging from 1.967 (8) to 1.980 (8) Å. The N–Fe–N bite angles are similar to those previously reported [82.8 (3)–83.1 (3)°]. Each phen ligand is planar, with mean deviations of

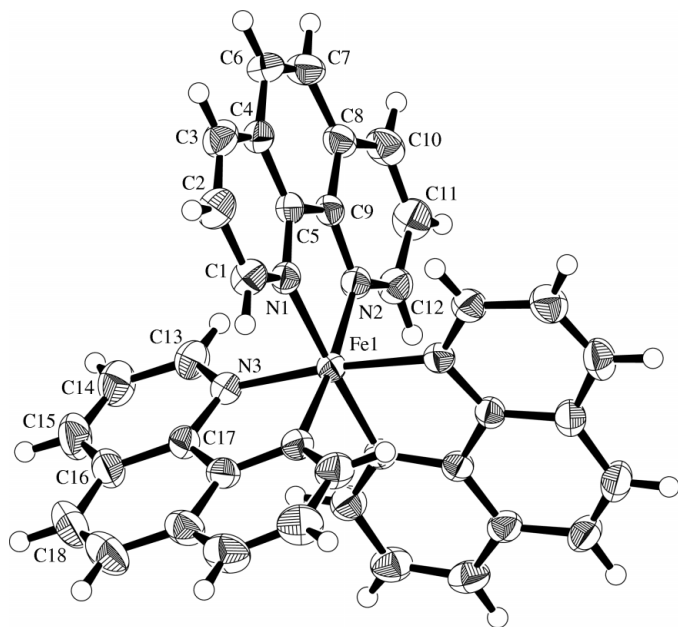


Figure 1
ORTEP (Johnson, 1976) drawing of the $[\text{Fe}(\text{phen})_3]^{3+}$ cation of (I). Displacement ellipsoids for non-H atoms are shown at the 50% probability level. Unlabeled atoms are related to labeled atoms by the symmetry operator $(-x, y, \frac{1}{2} - z)$.

0.036 [plane N1/C1–C12/N2] and 0.011 Å [plane N3/C13–C18/C13ⁱ–C18ⁱ/N3ⁱ; symmetry code as in Table 1].

The crystal structure is stabilized by partial stacking interactions between the phen ligands of neighboring cations [$\text{C4} \cdots \text{C7}^{\text{ii}}$ and $\text{C7} \cdots \text{C4}^{\text{ii}}$ = 3.383 (7), $\text{C14} \cdots \text{C15}^{\text{iii}}$ and $\text{C15} \cdots \text{C14}^{\text{iii}}$ = 3.589 (3) Å; symmetry codes: (ii) $-x, -y, -z$; (iii) $-\frac{1}{2} - x, \frac{1}{2} - y, -z$; Fig. 2] and by O–H \cdots O hydrogen-bonding interactions between the disordered water molecule (lying on a twofold axis) and nitrate anions (Table 2 and Fig. 3). One nitrate anion lies on a twofold rotation axis.

Experimental

Brown prismatic crystals of the title compound were obtained at room temperature by the slow evaporation of a DMF solution of 1,10-phenanthroline, iron(III) nitrate nonahydrate and malonic acid (molar ratio 1:1:1).

Crystal data

$[\text{Fe}(\text{C}_{12}\text{H}_8\text{N}_2)_3](\text{NO}_3)_3 \cdot \text{H}_2\text{O}$
 $M_r = 800.50$
 Monoclinic, $C2/c$
 $a = 10.769$ (8) Å
 $b = 24.58$ (2) Å
 $c = 13.274$ (12) Å
 $\beta = 103.00$ (3)°
 $V = 3423$ (5) Å³
 $Z = 4$

$D_x = 1.553$ Mg m⁻³
 Mo $K\alpha$ radiation
 Cell parameters from 12 716 reflections
 $\theta = 3.1$ – 27.4 °
 $\mu = 0.52$ mm⁻¹
 $T = 296.1$ K
 Block, brown
 $0.20 \times 0.20 \times 0.20$ mm

Data collection

Rigaku R-AXIS RAPID diffractometer
 ω scans
 Absorption correction: multi-scan (Higashi, 1995).
 $T_{\text{min}} = 0.732$, $T_{\text{max}} = 0.904$
 16 262 measured reflections

3869 independent reflections
 2512 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.024$
 $\theta_{\text{max}} = 27.4$ °
 $h = -12 \rightarrow 13$
 $k = -31 \rightarrow 31$
 $l = -17 \rightarrow 17$

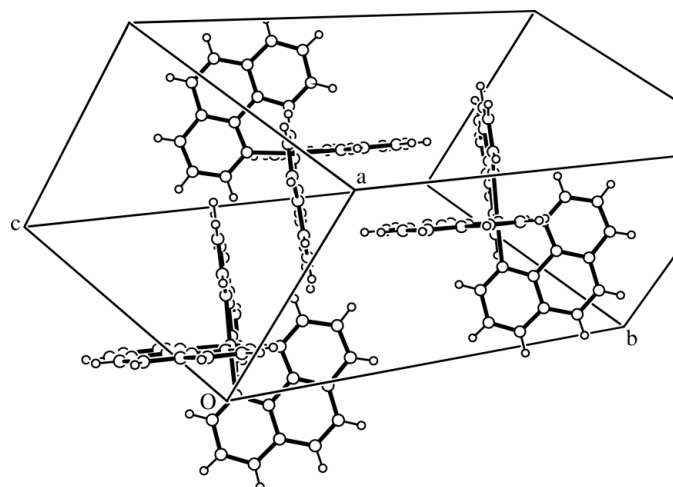


Figure 2
View of the crystal packing, showing stacking of phen ligands.

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.043$
 $wR(F^2) = 0.123$
 $S = 1.02$
 3869 reflections
 256 parameters

H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0811P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.71$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.49$ e Å⁻³

Table 1

Selected geometric parameters (Å, °).

Fe1–N1	1.988 (1)	Fe1–N3	1.968 (2)
Fe1–N2	1.971 (2)		
N1–Fe1–N1 ⁱ	177.73 (7)	N2–Fe1–N2 ⁱ	89.39 (7)
N1–Fe1–N2	82.43 (6)	N2–Fe1–N3	93.89 (7)
N1–Fe1–N2 ⁱ	95.95 (6)	N2–Fe1–N3 ⁱ	173.95 (6)
N1–Fe1–N3	89.53 (6)	N3–Fe1–N3 ⁱ	83.30 (7)
N1–Fe1–N3 ⁱ	92.17 (6)		

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

Table 2

Hydrogen-bonding geometry (Å, °).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
O6–H13 \cdots O1 ⁱ	0.87	2.02	2.657 (4)	129
O6–H13 \cdots O2 ⁱ	0.87	2.38	3.053 (5)	135
O6–H14 \cdots O4 ⁱⁱ	0.89	2.24	3.067 (8)	154

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

H atoms, except those of water molecules, were placed in calculated positions, with C–H = 0.93 Å, and refined as riding, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$. Those of water molecules were located in a difference Fourier map and were then refined as riding with $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{O})$; one of these H atoms is disordered over two positions across the twofold rotation axis on which the water molecule lies, while the other lies on the axis.

Data collection: *PROCESS-AUTO* (Rigaku/MSC & Rigaku Corporation, 2004); cell refinement: *PROCESS-AUTO*; data reduction: *CrystalStructure* (Rigaku/MSC & Rigaku Corporation, 2004); program(s) used to solve structure: *SIR97* (Altomare *et al.*, 1999); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997);

molecular graphics: *ORTEPII* (Johnson, 1976); software used to prepare material for publication: *CrystalStructure*.

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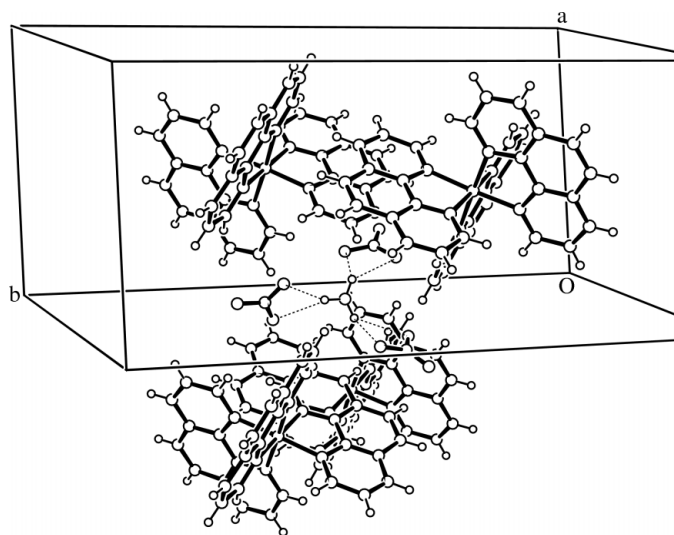


Figure 3
View of the crystal packing, showing hydrogen-bond interactions (dashed lines).

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