

Gareth R. Lewis,\*  
Peter Hubberstey and  
Claire L. SampsonSchool of Chemistry, University of Nottingham,  
Nottingham NG7 2RD, EnglandCorrespondence e-mail:  
gareth.lewis@nottingham.ac.uk

## Key indicators

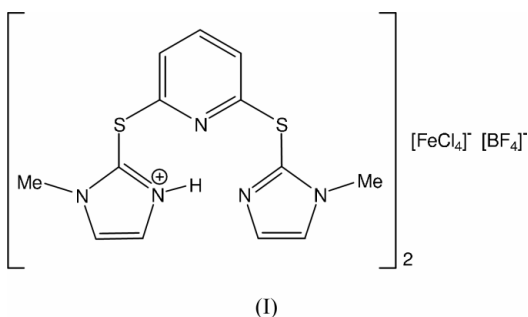
Single-crystal X-ray study  
 $T = 150$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.006$  Å  
 $R$  factor = 0.049  
 $wR$  factor = 0.095  
Data-to-parameter ratio = 18.5For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.Bis[2-(1-methyl-1*H*-imidazol-2-yl-3-*iosulfanyl*)-  
6-(1-methyl-1*H*-imidazol-2-ylsulfanyl)pyridine]  
tetrachloroferrate tetrafluoroborateThe cations of the title complex,  $[H\text{-}\{2,6\text{-}(\text{meth})_2\text{py}\}]_2\text{-}[\text{FeCl}_4][\text{BF}_4]$  or  $(\text{C}_{13}\text{H}_{14}\text{N}_5\text{S}_2)_2[\text{FeCl}_4]\text{BF}_4$ , form dimers of protonated 2,6-bis(1-methyl-1*H*-imidazol-2-ylsulfanyl)pyridine molecules, which are held together by moderate  $\text{N}-\text{H}\cdots\text{N}$  hydrogen bonds.

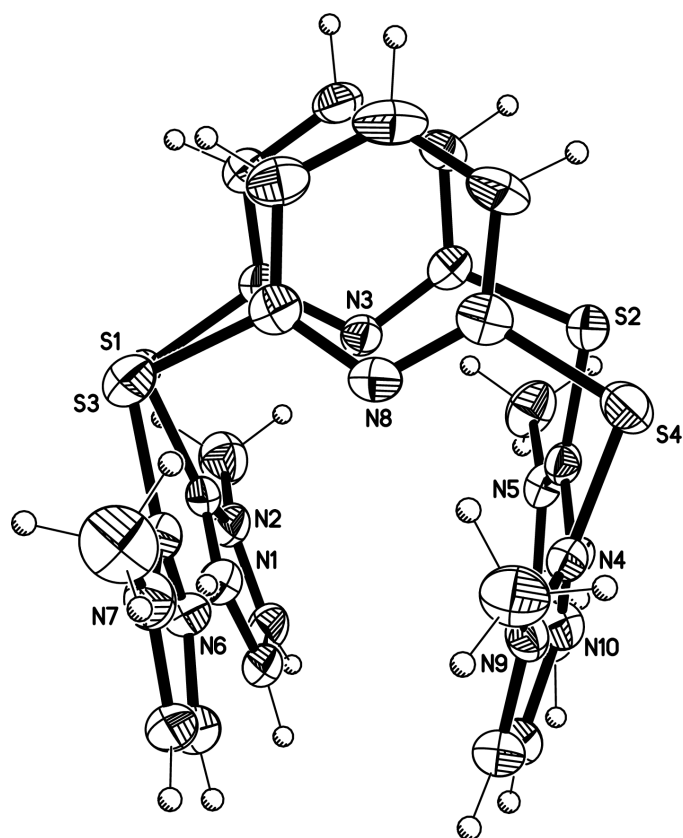
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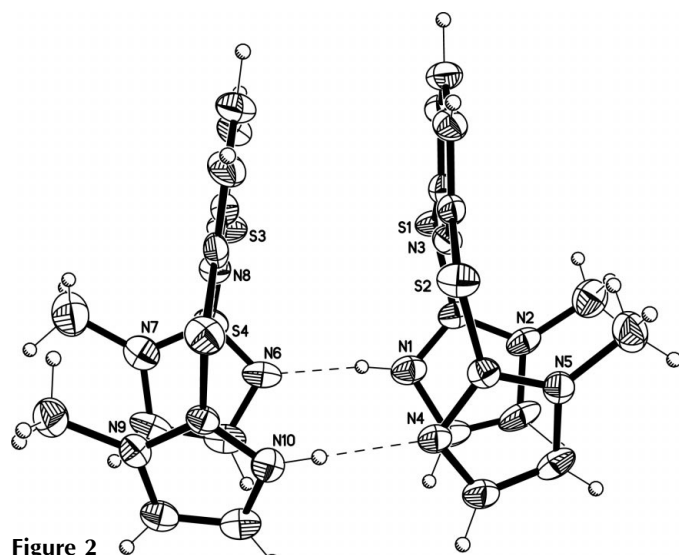
## Comment

Multidentate nitrogen-donor ligands are highly prevalent in transition-metal coordination chemistry and, with readily accessible synthetic routes, form an extremely large and diverse family of compounds. Consequently, these molecules may be employed in a wide variety of research areas with applications that include homogeneous catalysis, molecular magnetism, photochemistry and liquid crystals, in addition to fundamental research probing the chemistry of biological systems. While investigating the coordination chemistry of tripodal nitrogen-donor ligands, such as 2,6-bis(1-methyl-1*H*-imidazol-2-ylsulfanyl)pyridine [2,6-(meth)<sub>2</sub>py], the salt  $[H\text{-}\{2,6\text{-}(\text{meth})_2\text{py}\}]_2[\text{FeCl}_4][\text{BF}_4]$ , (I), was crystallized.The structure of (I) comprises two N-protonated 2-(1-methyl-1*H*-imidazol-2-yl-3-*iosulfanyl*)-6-(1-methyl-1*H*-imidazol-2-ylsulfanyl)pyridine molecules, one tetrachloroferrate and one tetrafluoroborate anion. On each organic moiety, one of the two methimazolyl heterocycles has a protonated imine N atom. Consequently, the cations form a dimer (Fig. 1), held together by moderately strong (Jeffery, 1997)  $\text{N}-\text{H}\cdots\text{N}$  hydrogen bonds. Within each cation, the angle at the S atom bridging between the pyridyl and methimazolyl rings prevents the two methimazolyl rings from being coplanar [the angle between the planes is  $26.2(2)^\circ$  in both molecules in the asymmetric unit]. The torsion of the methimazolyl rings about the C—S bonds appears to be relatively facile, with angles between the 1*H*-imidazol-2-ylsulfanyl and pyridyl ring planes of  $75.2(3)$ ,  $87.8(3)$ ,  $88.6(3)$  and  $93.4(3)^\circ$ . As shown in Fig. 2, the pyridyl rings of dimerized cations are slightly offset, the angle between the two planes being  $9.7(2)^\circ$ , with a closest

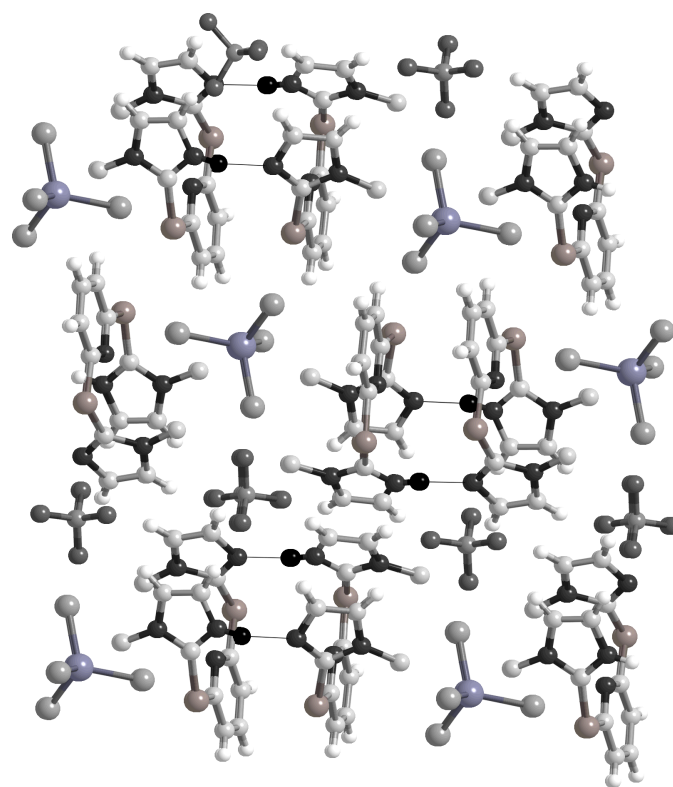


**Figure 1**  
Displacement ellipsoid plot of (I), with atoms represented at the 50% probability level. The anions have been omitted for clarity.

separation between these rings of 3.606 (3) Å (C7 and C20). Thus any  $\pi \cdots \pi$  interaction between the pyridyl rings is relatively weak, and is a consequence of the N—H $\cdots$ N hydrogen bonds between the cations.



**Figure 2**  
A second view of the cation dimer in the structure of (I), showing the offset face-to-face orientation of the pyridyl rings. Dashed lines indicate the N—H $\cdots$ N hydrogen bonds between the cations.



**Figure 3**  
The crystal structure of (I), with the N—H $\cdots$ N hydrogen bonds shown as thin black lines.

In the crystal structure of (I), the cation dimers are separated by both tetrafluoroborate and tetrachloroferrate anions (Fig. 3), with several moderate C—H $\cdots$ F and weak C—H $\cdots$ Cl hydrogen bonds between the ions (Table 2).

## Experimental

All reagents were purchased from Aldrich and used as received. All manipulations were carried out under an atmosphere of nitrogen, using dry deoxygenated solvents. A solution of methimazole (2.18 g, 19.1 mmol) in *N,N'*-dimethylformamide (DMF, 30 ml) was added to a suspension of sodium hydride (0.96 g, 24.0 mmol) in DMF (200 ml), and the mixture stirred at room temperature for 2 h. Subsequently, solid 2,6-dichloropyridine (1.46 g, 9.9 mmol) was added, and the mixture heated to reflux for 60 h. On cooling to room temperature, the mixture was poured over ice and the product extracted into CH<sub>2</sub>Cl<sub>2</sub>. The combined CH<sub>2</sub>Cl<sub>2</sub> fractions were dried over MgSO<sub>4</sub>, filtered and reduced to dryness *in vacuo*, yielding a yellow oil, which was recrystallized from hot ethanol, giving 2,6-bis(1-methyl-1*H*-imidazol-2-ylsulfanyl)pyridine as a pale-yellow solid (2.40 g, 7.9 mmol, 80%). Crystals of (I) suitable for X-ray diffraction studies were grown by allowing diethyl ether vapour to diffuse into an acetone solution containing iron(II) tetrafluoroborate hexahydrate and 2,6-bis(1-methyl-1*H*-imidazol-2-ylsulfanyl)pyridine over a period of 3 d.

## Crystal data

$2\text{C}_{13}\text{H}_{14}\text{N}_5\text{S}_2^+\cdot\text{FeCl}_4^-\cdot\text{BF}_4^-$   
 $M_r = 893.28$   
 Triclinic,  $P\bar{1}$   
 $a = 9.0531$  (10) Å  
 $b = 13.2213$  (14) Å  
 $c = 16.0249$  (18) Å  
 $\alpha = 82.51$  (2)°  
 $\beta = 82.04$  (3)°  
 $\gamma = 82.82$  (2)°  
 $V = 1872.2$  (4) Å<sup>3</sup>

$Z = 2$   
 $D_x = 1.585$  Mg m<sup>-3</sup>  
 Mo  $K\alpha$  radiation  
 Cell parameters from 391 reflections  
 $\theta = 1.9$ – $27.6$ °  
 $\mu = 0.97$  mm<sup>-1</sup>  
 $T = 150$  (2) K  
 Lath, yellow  
 $0.41 \times 0.14 \times 0.04$  mm

## Data collection

Bruker SMART 1000 CCD area-detector diffractometer  
 $\varphi$  and  $\omega$  scans  
 Absorption correction: multi-scan (*SAINT*; Bruker, 2001)  
 $T_{\min} = 0.693$ ,  $T_{\max} = 0.960$   
 10005 measured reflections

8426 independent reflections  
 4004 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.047$   
 $\theta_{\text{max}} = 29.0$ °  
 $h = -12 \rightarrow 10$   
 $k = -17 \rightarrow 17$   
 $l = -21 \rightarrow 15$

## Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.049$   
 $wR(F^2) = 0.095$   
 $S = 0.89$   
 8426 reflections  
 455 parameters

H atoms treated by a mixture of independent and constrained refinement  
 $w = 1/[\sigma^2(F_o^2) + (0.0259P)^2]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\text{max}} = 0.011$   
 $\Delta\rho_{\text{max}} = 0.53$  e Å<sup>-3</sup>  
 $\Delta\rho_{\text{min}} = -0.36$  e Å<sup>-3</sup>

Table 1

Selected geometric parameters (Å, °).

Fe1—Cl4	2.1864 (12)	Fe1—Cl3	2.1972 (11)
Fe1—Cl1	2.1902 (12)	Fe1—Cl2	2.1983 (11)
C4—S1—C5	101.14 (18)	C17—S3—C18	102.64 (18)
C10—S2—C9	101.97 (17)	C23—S4—C22	100.61 (18)

Table 2

Intermolecular hydrogen bonds (Å, °) in the crystal structure of (I).

$D-H\cdots A$	$D-H$	$D\cdots A$	$D-H\cdots A$
N1—H1N $\cdots$ N6	1.82	2.675 (3)	171
N10—H10N $\cdots$ N4	1.88	2.718 (3)	163
C1—H1A $\cdots$ F1	2.24	3.154 (3)	143
C25—H25A $\cdots$ F3	2.32	3.238 (3)	174
C26—H26C $\cdots$ F4	2.34	3.273 (3)	176
C2—H2A $\cdots$ F1	2.39	3.241 (3)	173
C24—H24A $\cdots$ F3	2.45	3.211 (3)	137
Cl3—H13B $\cdots$ Cl3	2.79	3.562 (3)	160

The H atoms of the cation were located in a difference map and constrained to idealized geometries. The methyl groups were allowed to rotate but not to tip.

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINT* (Bruker, 2001); data reduction: *SAINT* and *SHELXTL* (Bruker, 2001); 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* and *PLATON* (Spek, 2001).

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