

Gareth R. Owen\* and Mairi F. Haddow

School of Chemistry, Cantocks Close, University of Bristol, Bristol, BS8 1TS, UK

Correspondence e-mail:  
gareth.owen@bris.ac.uk

## Key indicators

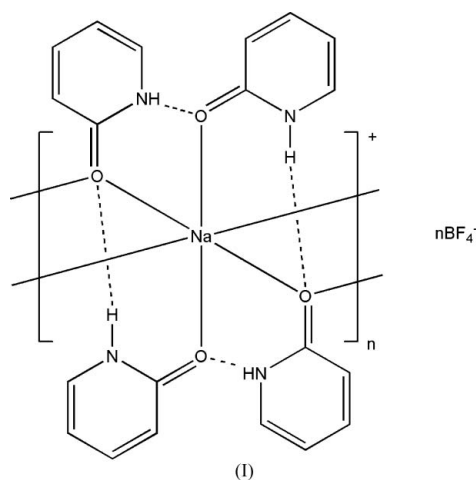
Single-crystal X-ray study  
 $T = 173$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.004$  Å  
 $R$  factor = 0.065  
 $wR$  factor = 0.147  
Data-to-parameter ratio = 15.5For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.*catena*-Poly[[[bis(2-pyridone- $\kappa$ O)sodium]-di- $\mu$ -2-pyridone- $\kappa^4$ O:O] tetrafluoroborate]

In the title compound,  $[\text{Na}(\text{C}_5\text{H}_5\text{NO})_4]\text{BF}_4$ , each  $\text{Na}^+$  ion, lying on an inversion centre, is octahedrally coordinated by four bridging and two terminal pyridone ligands, creating a cationic coordination polymer chain with tetrafluoroborate anions occupying the spaces between the chains.

Received 17 November 2006  
Accepted 28 November 2006

## Comment

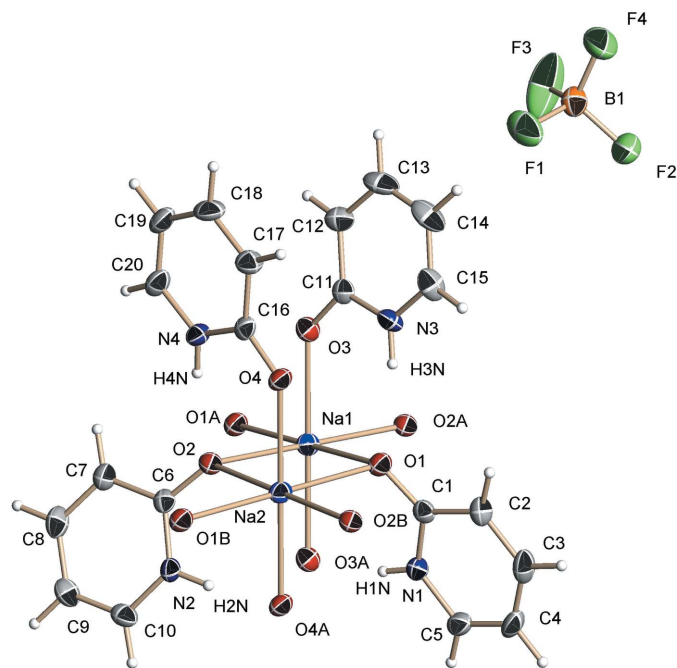
The title compound, (I), was prepared during the course of our investigations on the tautomerization of 2-pyridone and 2-mercaptopyridine rings in the preparation of new scorpionate ligands. We are currently investigating the effect of tautomerization on ligand coordination properties.



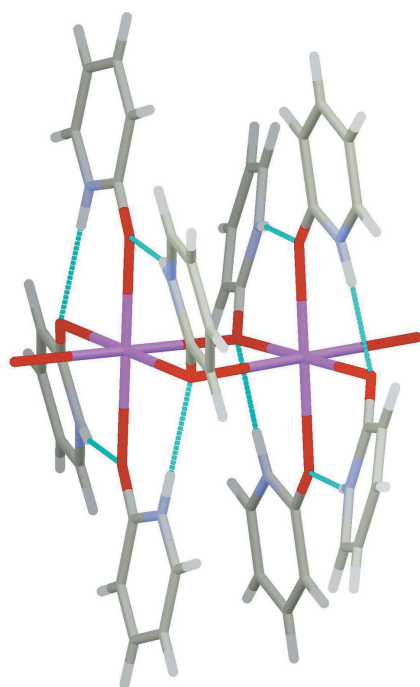
The compound forms a coordination polymer chain with two independent  $\text{Na}(\text{pyridone})_4^+$  units, each sodium sitting on an inversion centre. Each  $\text{Na}^+$  ion is connected to the sodium on either side by two bridging pyridone groups and is also coordinated by two terminal pyridone ligands, making each sodium octahedrally coordinated (Fig. 1 and Table 1). The pyridone ligands are arranged such that the amide H atom on each pyridone is hydrogen bonded to the oxygen of an adjacent pyridone ligand (Table 2), and these  $\text{Na}(\text{pyridone})_4^+$  groups stack together to form the coordination polymer (Fig. 2), with the  $\text{BF}_4^-$  anions occupying the spaces between the chains (Fig. 3). Some  $\pi$  stacking occurs between the 2-pyridone rings with minimum distances between the atoms in the rings of 3.125 (3) (C11–C16) and 3.284 (4) Å [C16–N1<sup>ii</sup>; symmetry code: (ii)  $2 - x, 1 - y, 1 - z$ ].

## Experimental

A round-bottomed flask was charged with  $\text{NaBF}_4$  (0.05 g, 0.45 mmol) and methanol (40 ml). The mixture was stirred until the salt had

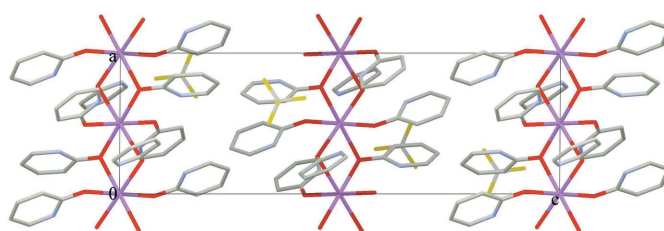


**Figure 1**  
The  $\text{Na}(\text{pyridone})_4^+$  and  $\text{BF}_4^-$  ions with the atom-labelling scheme. Displacement ellipsoids are drawn at the 50% probability level with H atoms represented as spheres of arbitrary size. [Symmetry codes: (A)  $1 - x, 1 - y, 1 - z$ ; (B)  $2 - x, 1 - y, 1 - z$ .



**Figure 2**  
Hydrogen-bonded (dashed lines) units are stacked to form a coordination polymer structure.

dissolved. 2-Hydroxypyridine (0.17 g, 1.80 mmol) was subsequently added to the mixture, which was left to stir overnight. The solvent was removed under reduced pressure and the residue was dissolved in the minimum amount of THF. The resulting solution was left to stand at 273 K for 48 h, resulting in the formation of colourless needles



**Figure 3**  
Packing diagram, viewed along the  $b$  axis. H atoms have been omitted.

(0.21 g, 0.43 mmol, 95%). Calcd for  $\text{C}_{20}\text{H}_{20}\text{N}_4\text{O}_4\text{F}_4\text{BNa}$ : C, 49.00; H, 4.11; N, 11.43. Found: C, 49.00; H, 4.14; N, 11.77.

**Crystal data**

$[\text{Na}(\text{C}_5\text{H}_5\text{NO})_4]\text{BF}_4$   
 $M_r = 490.20$   
Monoclinic,  $P2_1/n$   
 $a = 7.2259(12) \text{ \AA}$   
 $b = 13.521(2) \text{ \AA}$   
 $c = 22.655(4) \text{ \AA}$   
 $\beta = 90.322(4)^\circ$   
 $V = 2213.5(6) \text{ \AA}^3$

$Z = 4$   
 $D_x = 1.471 \text{ Mg m}^{-3}$   
Mo  $K\alpha$  radiation  
 $\mu = 0.14 \text{ mm}^{-1}$   
 $T = 173(2) \text{ K}$   
Needle, colourless  
 $0.42 \times 0.06 \times 0.06 \text{ mm}$

**Data collection**

Bruker SMART CCD area-detector diffractometer  
 $\varphi$  and  $\omega$  scans  
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)  
 $T_{\text{min}} = 0.849, T_{\text{max}} = 0.992$

22997 measured reflections  
5053 independent reflections  
3175 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.076$   
 $\theta_{\text{max}} = 27.5^\circ$

**Refinement**

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.065$   
 $wR(F^2) = 0.147$   
 $S = 1.03$   
5053 reflections  
326 parameters  
H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0576P)^2 + 1.6958P]$   
where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\text{max}} < 0.001$   
 $\Delta\rho_{\text{max}} = 0.31 \text{ e \AA}^{-3}$   
 $\Delta\rho_{\text{min}} = -0.26 \text{ e \AA}^{-3}$

**Table 1**

Selected geometric parameters ( $\text{\AA}, ^\circ$ ).

Na1—O1	2.4099 (18)	Na2—O1	2.5422 (19)
Na1—O2	2.4940 (19)	Na2—O2	2.4043 (18)
Na1—O3	2.3122 (19)	Na2—O4	2.3172 (19)
Na1—O1—Na2	93.66 (6)	Na1—O2—Na2	95.04 (6)

**Table 2**

Hydrogen-bond geometry ( $\text{\AA}, ^\circ$ ).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
N1—H1N $\cdots$ O3 <sup>i</sup>	0.82 (3)	1.97 (3)	2.772 (3)	168 (3)
N2—H2N $\cdots$ O4 <sup>ii</sup>	0.88 (3)	1.92 (3)	2.776 (3)	163 (3)
N3—H3N $\cdots$ O1	0.85 (3)	2.00 (3)	2.838 (3)	169 (3)
N4—H4N $\cdots$ O2	0.87 (3)	1.96 (3)	2.827 (3)	169 (3)

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

H atoms were treated as riding, with C–H = 0.95 Å and  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$  for aromatic H atoms. Amine protons were refined isotropically.

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINTE* (Bruker, 1998); data reduction: *SAINTE*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 2001); software used to prepare material for publication: *SHELXTL*.

The authors thank the Ramsay Trust (GO) and EPSRC (MH) for their financial support.

## References

- Bruker (1998). *SMART* (Version 5.054) and *SAINTE* (Version 6.02). Bruker AXS Inc., Madison, Wisconsin, USA.
- Bruker (2001). *SHELXTL*. Version 6.12. Bruker AXS Inc., Madison, Wisconsin, USA.
- Sheldrick, G. M. (1996). *SADABS*. University of Göttingen, Germany.
- Sheldrick, G. M. (1997). *SHELXS97* and *SHELXL97*. University of Göttingen, Germany.