# metal-organic papers

Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

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#### **Key indicators**

Single-crystal X-ray study T = 173 KMean  $\sigma$ (C–C) = 0.018 Å R factor = 0.046 wR factor = 0.102 Data-to-parameter ratio = 22.7

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

# A bis[pentaiodobismuthate(III)] salt of 4-hydroxypyridinium

The crystal structure of tetra(4-hydroxypyridinium) di- $\mu$ iodo-bis[tetraiodobismuthate(III)] 4-pyridone tetrasolvate,  $(C_5H_6NO)_4[Bi_2I_{10}]\cdot 4C_5H_5NO$ , has been determined. The structure contains discrete centrosymmetric  $[Bi_2I_{10}]^{4-}$  anions enclosed in a hydrogen-bonded array of 4-hydroxypyridinium cations and 4-pyridone molecules. Received 1 February 2002 Accepted 15 February 2002 Online 15 March 2002

## Comment

The chemistry of iodoantimonate and iodobismuthate anions has been an area of some interest (Fisher & Norman, 1994). Iodobismuthate anions exhibit considerable structural variety and examples of both polymeric (Carmalt *et al.*, 1995; Geiser *et al.*, 1990; Krautscheid, 1995; Rogers *et al.*, 1992) and oligomeric (Clegg *et al.*, 1991; Eickmeier *et al.*, 1999; Krautscheid, 1994; Kubiak & Ejsmont, 1999) forms are known. Of the numerous counter-cations present, we note, in relation to this study, a dinuclear iodobismuthate with a hydrogen-bonded cation, namely  $[2,2'-Hbipy]_4[Bi_2I_{10}]$  (Bowmaker et al., 1998). Herein we report the structure of a dinuclear iodobismuthate, (I), in which the 4-hydroxypyridinium cations form a hydrogen-bonded network with neutral 4-pyridone molecules.



One of the products of the reaction of 4-pyridone and diiodophenylbismuth(III) in tetrahydrofuran (THF) is the title compound, a salt containing a dimeric periodobismuth tetraanion and a 1:1 mixture of hydroxypyridinium counterions and 4-pyridone starting material (Fig. 1). The hydrogen bonding shown in Fig. 1, combined with interactions between H2A and O4(1-x, 2-y, 2-z), gives rise to a hydrogen-bonded set of close-packed ribbons of 4-hydroxypyridinium cations and 4pyridone molecules (Fig. 2) into which the anions are incorporated. Hydrogen-bond N···O and O···O distances within the ribbons range from 2.518 (11) to 2.715 (12) Å. Much weaker interactions are seen between the ribbons and the iodo ligands of the bismuthate anions; the shortest N···I hydrogenbonding contact is 3.627 (9) Å (Table 2).

### **Experimental**

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Crystals of the title compound were obtained as a minor product from the reaction between  $BiI_2Ph$  (0.100 g, 0.185 mmol) and 4-pyridone (0.020 g, 0.21 mmol), both dissolved in THF, to which an overlayer of



#### Figure 1

The molecular structure of the title compound, showing 50% probability displacement ellipsoids. Hydrogen bonds are shown as dashed lines.

hexane was added. Solvent diffusion over a period of some days at room temperature afforded a small quantity of red crystals.

#### Crystal data

$(C_5H_6NO)_4[Bi_2I_{10}]\cdot 4C_5H_5NO$	Z = 1
$M_r = 2451.80$	$D_x = 2.646 \text{ Mg m}^{-3}$
Triclinic, P1	Mo $K\alpha$ radiation
a = 10.366 (3) Å	Cell parameters from 5958
b = 12.362 (3) Å	reflections
c = 13.504(5) Å	$\theta = 2-25^{\circ}$
$\alpha = 116.212 \ (13)^{\circ}$	$\mu = 10.78 \text{ mm}^{-1}$
$\beta = 95.05 \ (2)^{\circ}$	T = 173 (2) K
$\gamma = 92.914 \ (16)^{\circ}$	Lath, red
$V = 1538.6 (8) \text{ Å}^3$	$0.15 \times 0.05 \times 0.02 \text{ mm}$

#### Data collection

6960 independent reflections
4015 reflections with $I > 2\sigma(I)$
$R_{\rm int} = 0.061$
$\theta_{\rm max} = 27.5^{\circ}$
$h = -13 \rightarrow 12$
$k = -16 \rightarrow 16$
$l = -17 \rightarrow 17$



#### Figure 2

Packing diagram, viewed down the a axis, showing hydrogen-bonded ribbons of 4-pyridone and 4-hydroxypyridinium. The bismuth-containing anions have been omitted for clarity.

Refinement	
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Refinement on $F^2$	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.046$	$w = 1/[\sigma^2(F_o^2) + (0.0377P)^2]$
$wR(F^2) = 0.102$	where $P = (F_o^2 + 2F_c^2)/3$
S = 0.90	$(\Delta/\sigma)_{\rm max} = 0.001$
6960 reflections	$\Delta \rho_{\rm max} = 1.87 \text{ e } \text{\AA}^{-3}$
307 parameters	$\Delta \rho_{\rm min} = -2.67 \text{ e } \text{\AA}^{-3}$

#### Table 1

Selected geometric parameters (Å, °).

Bi1-I1	2.9657 (12)	O4-C18	1.279 (13)
Bi1-I3	3.0082 (12)	C1-C2	1.379 (17)
Bi1-I4	3.0502 (13)	C2-C3	1.391 (16)
Bi1-I2	3.1056 (13)	C3-C4	1.390 (17)
Bi1-I5	3.2099 (12)	C4-C5	1.355 (17)
Bi1-I5 <sup>i</sup>	3.2197 (12)	C6-C7	1.355 (15)
I5-Bi1 <sup>i</sup>	3.2197 (12)	C7-C8	1.408 (16)
N1-C1	1.324 (15)	C8-C9	1.416 (16)
N1-C5	1.329 (15)	C9-C10	1.341 (16)
N2-C10	1.343 (14)	C11-C12	1.356 (16)
N2-C6	1.374 (15)	C12-C13	1.377 (16)
N3-C11	1.335 (16)	C13-C14	1.427 (16)
N3-C15	1.345 (15)	C14-C15	1.355 (16)
N4-C20	1.340 (14)	C16-C17	1.357 (16)
N4-C16	1.341 (15)	C17-C18	1.439 (15)
O1-C3	1.329 (14)	C18-C19	1.424 (16)
O2-C8	1.311 (13)	C19-C20	1.359 (16)
O3-C13	1.317 (13)		
I1-Bi1-I3	95.43 (4)	I4-Bi1-I5	92.67 (3)
I1-Bi1-I4	91.10 (3)	I2-Bi1-I5	90.28 (3)
I3-Bi1-I4	88.77 (4)	$I1-Bi1-I5^{i}$	86.95 (4)
I1-Bi1-I2	85.81 (3)	I3-Bi1-I5 <sup>i</sup>	177.04 (3)
I3-Bi1-I2	93.51 (4)	$I4-Bi1-I5^{i}$	89.43 (4)
I4-Bi1-I2	176.31 (3)	I2-Bi1-I5 <sup>i</sup>	88.40 (4)
I1-Bi1-I5	174.51 (3)	I5-Bi1-I5 <sup>i</sup>	89.09 (4)
I3-Bi1-I5	88.65 (4)	Bi1-I5-Bi1 <sup>i</sup>	90.91 (4)

Symmetry code: (i) 1 - x, 1 - y, 1 - z.

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

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
$N1 - H1A \cdots O4$	0.88	1.78	2.646 (14)	168
$N2-H2A\cdots O4^{i}$	0.88	1.87	2.715 (12)	160
$N3-H3A\cdots I2^{ii}$	0.88	3.29	3.870 (11)	126
$N3-H3A\cdots I4^{iii}$	0.88	2.87	3.633 (10)	147
$N4-H4A\cdots I1^{iv}$	0.88	3.25	3.898 (11)	133
N4-H4 $A$ ···I2 <sup>iv</sup>	0.88	2.89	3.627 (9)	144
$O1-H1\cdots O2$	0.84	1.75	2.567 (12)	164
O3−H3···O2	0.84	1.70	2.518 (11)	163

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

The hydroxyl H atoms were assigned to the rings in which the bond lengths show the least deviation from aromaticity (Table 1). These H atoms could not be located in a difference map and were assigned fixed positions in the plane of the pyridinium ring appropriate for hydrogen bonding to nearby acceptors, using the AFIX 83 instruction in *SHELXTL* (Bruker, 1998);  $U_{\rm iso}({\rm H}) = 1.5U_{\rm eq}({\rm O})$ . All other H atoms were constrained to ideal geometries and assigned isotropic displacement parameters 1.2 times those of their parent C atoms. The highest residual electron-density peak is found 0.92 Å from Bi1. Other residual electron-density peaks with values in the range 1.17– 1.58 e Å<sup>-3</sup> are also found near the Bi1 atom.

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINT* (Bruker, 1998); data reduction: *SHELXTL* (Bruker, 1998);

# metal-organic papers

program(s) used to solve structure: *SHELXTL*; program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

We would like to thank the Cambridge Crystallographic Data Centre for project studentship funding (JS).

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