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Key indicators

Single-crystal X-ray study
T = 291 K
Mean $\sigma(\text{C}-\text{C}) = 0.010 \text{ \AA}$
R factor = 0.036
wR factor = 0.100
Data-to-parameter ratio = 26.1

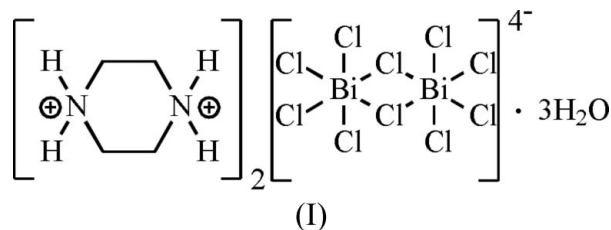
For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

Bis(piperazinium) di- μ -chloro-bis[tetrachloro-bismuthate(III)] trihydrate

The title salt, $(\text{C}_4\text{H}_{12}\text{N}_2)_2[\text{Bi}_2\text{Cl}_{10}]\cdot 3\text{H}_2\text{O}$, was prepared by reaction of bismuth trichloride and piperazine in a hydrochloric acid medium. It consists of piperazinium dications and dimeric $[\text{Bi}_2\text{Cl}_{10}]^{4-}$ tetraanions, the latter with twofold rotation symmetry, together with water of crystallization. Hydrogen-bond interactions, including $\text{N}-\text{H}\cdots\text{O}$, $\text{N}-\text{H}\cdots\text{Cl}$ and $\text{O}-\text{H}\cdots\text{Cl}$ interactions, stabilize the crystal structure.

Comment

There is increasing interest in halobismuthate(III) compounds, due to their anti-ulcer activity (Turel *et al.*, 1998) and their unique optical and electronic properties, including nonlinear optical activity, luminescence and semiconductivity (Goforth *et al.*, 2004). We report here the crystal structure of the title organic-inorganic hybrid complex, (I), of di- μ -chlorooctachlorodibismuthate with piperazinium.



Complex (I) consists of piperazinium(2+) dications and dimeric $[\text{Bi}_2\text{Cl}_{10}]^{4-}$ tetraanions (Figs. 1 and 2), together with water molecules of crystallization. The $[\text{Bi}_2\text{Cl}_{10}]^{4-}$ anion is located on a crystallographic C_2 symmetric axis, with the two $[\text{BiCl}_6]$ octahedra sharing one edge formed by two bridging Cl^- anions. The piperazinium cation has a chair-like conformation.

There are complicated hydrogen-bonding interactions among the $[\text{Bi}_2\text{Cl}_{10}]^{4-}$ anions, piperazinium cations and water molecules of crystallization. In particular, the $\text{N}-\text{H}\cdots\text{Cl}$ hydrogen bonds between each of the $[\text{Bi}_2\text{Cl}_{10}]^{4-}$ anions and its four neighbouring piperazinium cations result in interesting chains, formed by two parallel rows of piperazinium cations sandwiched between two rows of $[\text{Bi}_2\text{Cl}_{10}]^{4-}$ anions, along the c axis in the crystal structure (Fig. 3). These chains are then linked by hydrogen bonds involving the water molecules, forming an extended three-dimensional structure.

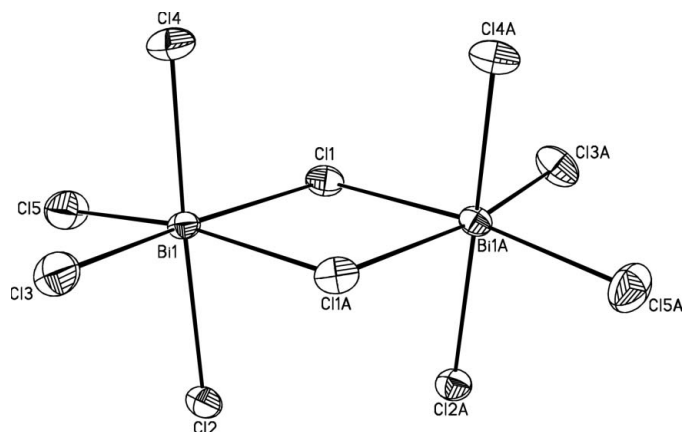
Experimental

Colourless crystals of (I) suitable for X-ray diffraction studies were obtained by slow evaporation for about 4 d at room temperature of an aqueous solution formed by mixing aqueous solutions of BiCl_3

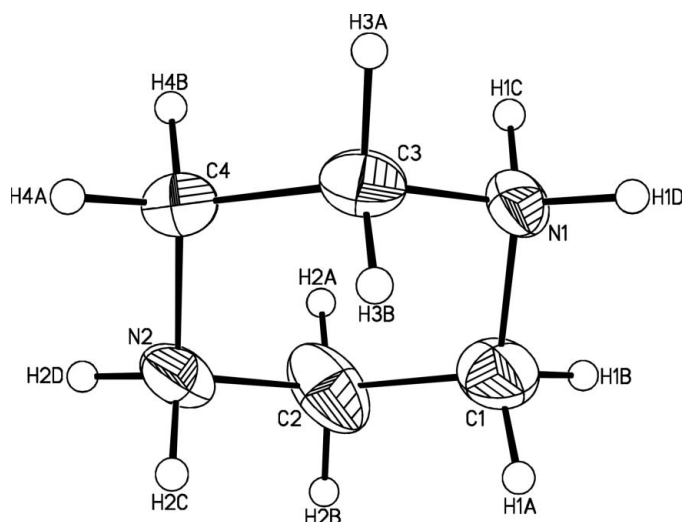
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Figure 1

A drawing of the $[\text{Bi}_2\text{Cl}_{10}]^{4-}$ anion of (I), with the atomic numbering scheme. Displacement ellipsoids are drawn at the 50% probability level. [Symmetry code: (A) $1 - x, y, \frac{3}{2} - z$.]


Figure 2

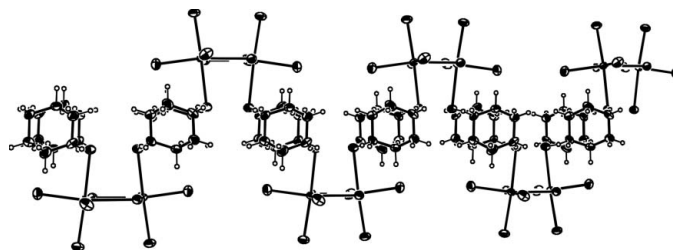
A drawing of the piperazium cation of (I), with the atomic numbering scheme. Displacement ellipsoids are drawn at the 50% probability level.

(1.24 mmol, dissolved in 2 ml concentrated HCl) and piperazine hexahydrate (4.94 mmol, dissolved in 15 ml H_2O). Spectroscopic analysis: IR (KBr, ν , cm^{-1}): 3546 (*m*, O—H), 3466 (*m*, O—H), 3124 (*m*, N—H), 3064 (*m*, N—H), 2810 (*m*, C—H), 2520 (*w*), 2427 (*w*), 1594 (*vs*, H_2O), 1444 (*vs*, CH_2), 1417 (*m*), 1407 (*m*), 1315 (*m*), 1201 (*w*), 1080 (*s*), 1054 (*s*), 1040 (*m*), 999 (*m*), 941 (*s*), 863 (*m*), 809 (*w*), 682 (*m*); UV (in DMSO): 256, 314 nm.

Crystal data

$(\text{C}_4\text{H}_{12}\text{N}_2)[\text{Bi}_2\text{Cl}_{10}] \cdot 3\text{H}_2\text{O}$
 $M_r = 1002.82$
 Monoclinic, $C2/c$
 $a = 14.3223$ (14) Å
 $b = 15.7868$ (14) Å
 $c = 13.5226$ (13) Å
 $\beta = 115.067$ (2)°
 $V = 2769.5$ (5) Å³
 $Z = 4$

$D_x = 2.405$ Mg m⁻³
 Mo $K\alpha$ radiation
 Cell parameters from 3992 reflections
 $\theta = 2.9$ – 28.2 °
 $\mu = 13.67$ mm⁻¹
 $T = 291$ (2) K
 Block, colourless
 $0.30 \times 0.20 \times 0.15$ mm


Figure 3

A view showing the chain formed by the $[\text{Bi}_2\text{Cl}_{10}]^{4-}$ anions and piperazium cations via N—H...Cl hydrogen bonds.

Data collection

Bruker APEX CCD area-detector diffractometer
 φ and ω scans
 Absorption correction: multi-scan (SADABS; Sheldrick 1996)
 $T_{\text{min}} = 0.053$, $T_{\text{max}} = 0.132$
 8270 measured reflections

3240 independent reflections
 2631 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.067$
 $\theta_{\text{max}} = 28.3$ °
 $h = -17 \rightarrow 18$
 $k = -13 \rightarrow 20$
 $l = -17 \rightarrow 17$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.036$
 $wR(F^2) = 0.100$
 $S = 1.06$
 3240 reflections
 124 parameters
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0408P)^2 + 9.2258P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.001$
 $\Delta\rho_{\text{max}} = 2.20 \text{ e } \text{Å}^{-3}$
 $\Delta\rho_{\text{min}} = -3.00 \text{ e } \text{Å}^{-3}$
 Extinction correction: SHELXL97 (Sheldrick, 1997)
 Extinction coefficient: 0.00346 (12)

Table 1

Selected geometric parameters (Å, °).

Bi—Cl3	2.5806 (18)	N1—C3	1.479 (7)
Bi—Cl5	2.5828 (18)	N1—C1	1.481 (8)
Bi—Cl4	2.6344 (16)	N2—C4	1.471 (8)
Bi—Cl2	2.8034 (16)	N2—C2	1.475 (8)
Bi—Cl1	2.8493 (19)	C1—C2	1.496 (9)
Bi—Cl1 ⁱ	2.8652 (18)	C3—C4	1.488 (8)
Cl1—Bi ⁱ	2.8652 (18)		
Cl3—Bi—Cl5	94.08 (7)	Cl5—Bi—Cl1 ⁱ	171.53 (5)
Cl3—Bi—Cl4	93.53 (5)	Cl4—Bi—Cl1 ⁱ	98.72 (5)
Cl5—Bi—Cl4	89.74 (6)	Cl2—Bi—Cl1 ⁱ	83.54 (4)
Cl3—Bi—Cl2	92.02 (5)	Cl1—Bi—Cl1 ⁱ	83.75 (5)
Cl5—Bi—Cl2	88.00 (6)	Bi—Cl1—Bi ⁱ	96.25 (5)
Cl4—Bi—Cl2	174.15 (4)	C3—N1—C1	109.4 (5)
Cl3—Bi—Cl1	169.16 (6)	C4—N2—C2	111.9 (5)
Cl5—Bi—Cl1	96.71 (6)	N1—C1—C2	112.1 (6)
Cl4—Bi—Cl1	85.54 (4)	N2—C2—C1	110.4 (6)
Cl2—Bi—Cl1	89.35 (4)	N1—C3—C4	111.8 (5)
Cl3—Bi—Cl1 ⁱ	85.72 (7)	N2—C4—C3	110.6 (5)
Cl3—Bi—Cl1—Bi ⁱ	13.2 (3)	C4—N2—C2—C1	55.0 (8)
Cl5—Bi—Cl1—Bi ⁱ	−172.10 (5)	N1—C1—C2—N2	−55.7 (8)
Cl4—Bi—Cl1—Bi ⁱ	98.69 (5)	C1—N1—C3—C4	−56.6 (6)
Cl2—Bi—Cl1—Bi ⁱ	−84.18 (4)	C2—N2—C4—C3	−55.6 (7)
Cl1 ⁱ —Bi—Cl1—Bi ⁱ	−0.61 (5)	N1—C3—C4—N2	56.7 (7)
C3—N1—C1—C2	56.2 (7)		

Symmetry code: (i) $-x + 1, y, -z + \frac{3}{2}$.

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>
N1—H1NB···O1	0.90	1.92	2.814 (6)	173
N1—H1NA···Cl2	0.90	2.53	3.291 (5)	143
N2—H2NA···O2	0.90	2.16	3.017 (5)	160
N2—H2NB···Cl2 ⁱⁱ	0.90	2.43	3.278 (5)	157
O1—H1···Cl4 ⁱⁱⁱ	0.85	2.48	3.318 (4)	170
O1—H2···Cl3 ⁱ	0.85	2.67	3.407 (5)	147
O2—H3···Cl5 ^{iv}	0.85	2.49	3.301 (5)	159

Symmetry codes: (i) $-x + 1, y, -z + \frac{3}{2}$; (ii) $-x + 1, -y + 2, -z + 2$; (iii) $-x + \frac{3}{2}, -y + \frac{3}{2}, -z + 2$; (iv) $x + \frac{1}{2}, y + \frac{1}{2}, z$.

H atoms attached to N and O atoms were located in a difference Fourier map and other H atoms were positioned geometrically. H atoms attached to N and C atoms were refined using a riding model, with C—H = 0.97 Å and N—H = 0.90 Å, and those attached to O atoms were refined in fixed positions, with O—H = 0.85 Å. For all H atoms, $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{parent atom})$. The locations of the maximum and

minimum difference density peaks are 0.95 and 0.85 Å, respectively, from Bi.

Data collection: *SMART* (Bruker, 2003); cell refinement: *SAINTE* (Bruker, 2003); 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, 2003); software used to prepare material for publication: *SHELXTL*.

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