

Acta Crystallographica Section E

Structure Reports

Online

ISSN 1600-5368

Poly[aqua- μ -bromido-(μ_2 -5-methyl-pyrazine-2-carboxylato- $\kappa^4N^1,O^2:O^2,O^2'$)lead(II)]

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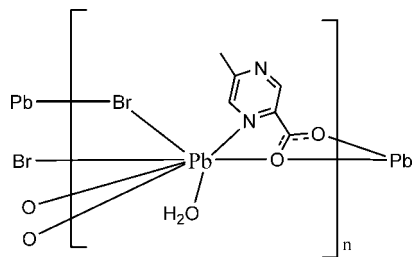
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Received 22 July 2012; accepted 27 July 2012

 Key indicators: single-crystal X-ray study; $T = 296$ K; mean $\sigma(C-C) = 0.010$ Å; R factor = 0.033; wR factor = 0.091; data-to-parameter ratio = 15.9.

In the title coordination polymer, $[PbBr(C_6H_5N_2O_2)(H_2O)]_n$, the Pb^{II} atom is coordinated by one pyrazine N atom, two bridging Br atoms, a water molecule and three carboxylate O atoms. Bridging by the two anions generates a layer structure parallel to (001); the layers are linked by $O-H \cdots N$ and $O-H \cdots Br$ hydrogen bonds, forming a three-dimensional network. The lone pair is stereochemically active, resulting in a Ψ -dodecahedral coordination environment for Pb^{II} .

Related literature

 For background, see: Ding *et al.* (2009).


Experimental

Crystal data

 $[PbBr(C_6H_5N_2O_2)(H_2O)]$
 $M_r = 442.24$

 Monoclinic, $P2_1/c$
 $a = 7.5493$ (10) Å
 $b = 6.6775$ (9) Å
 $c = 19.335$ (3) Å
 $\beta = 92.884$ (2)°
 $V = 973.5$ (2) Å³
 $Z = 4$
 Mo $K\alpha$ radiation
 $\mu = 21.41$ mm⁻¹
 $T = 296$ K
 $0.15 \times 0.14 \times 0.13$ mm

Data collection

 Bruker SMART APEX CCD area-detector diffractometer
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
 $T_{min} = 0.142$, $T_{max} = 0.167$

 5123 measured reflections
 1904 independent reflections
 1747 reflections with $I > 2\sigma(I)$
 $R_{int} = 0.039$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.033$
 $wR(F^2) = 0.091$
 $S = 1.05$
 1904 reflections

 120 parameters
 H-atom parameters constrained
 $\Delta\rho_{max} = 2.26$ e Å⁻³
 $\Delta\rho_{min} = -2.20$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
$O3-H3A \cdots N2^i$	0.85	1.97	2.816 (9)	173
$O3-H3B \cdots Br1^{ii}$	0.85	2.56	3.378 (6)	161

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

Data collection: APEX2 (Bruker, 2007); cell refinement: SAINT (Bruker, 2007); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: DIAMOND (Brandenburg, 1999); software used to prepare material for publication: publCIF (Westrip, 2010).

This present work was supported by the Tianjin Educational Committee (20090504) and Tianjin Normal University (1E0402B).

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: NG5284).

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supporting information

Acta Cryst. (2012). E68, m1139 [doi:10.1107/S1600536812033776]

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S1. Experimental

Lead(II) bromide (73.4 mg, 0.2 mmol), 5-pyrazine-2-carboxylic acid (27.6 mg, 0.2 mmol) and water (15 ml) were sealed in a 25 ml Teflon-lined steel vessel. The mixture was heated to 393 K for 5 days. The autoclave was cooled to room temperature at a rate of 10 K h⁻¹. Yellow block-shaped crystals were obtained in 60% yield based on Pb.

S2. Refinement

H atoms were placed in calculated positions as riding atoms attached to non-riding atoms with O—H es of 0.85 Å and C—H 0.93 to 0.96 Å, and with $U_{\text{iso}}(\text{H}) = 1.2$ to $1.5U_{\text{eq}}(\text{O,C})$. The final difference Fourier map had a peak in the vicinity of Pb and a hole in the vicinity of the same atom.

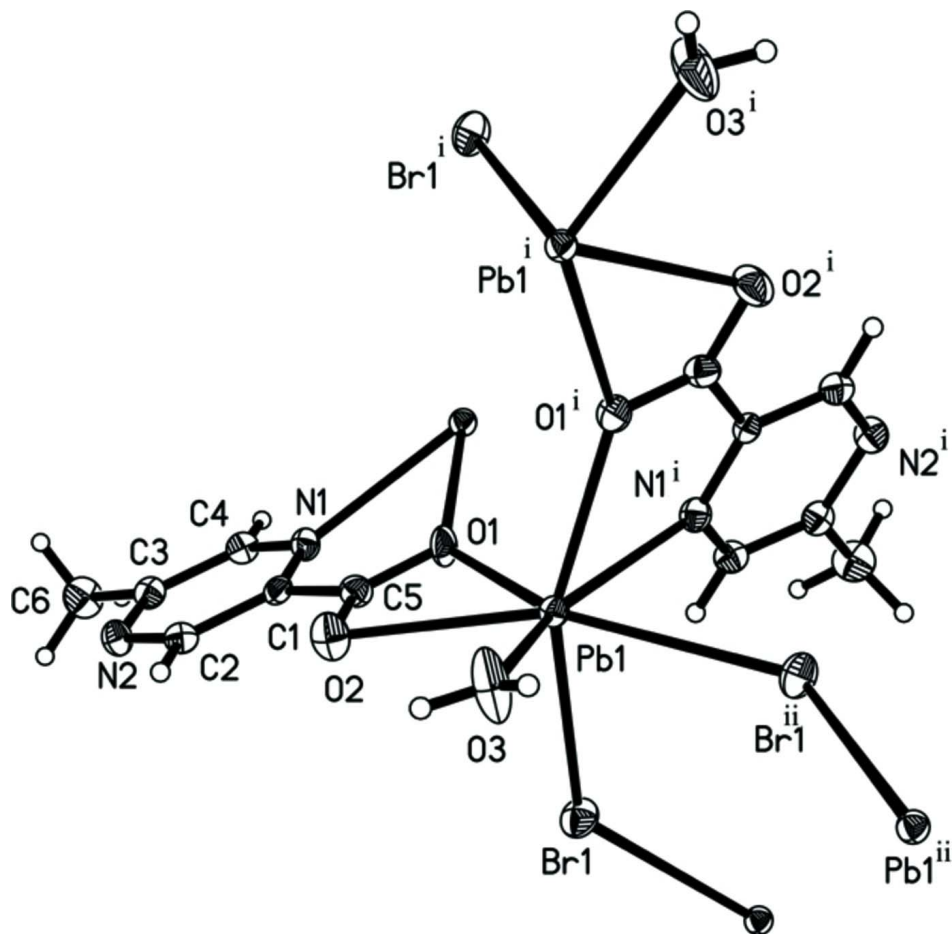


Figure 1

ORTEP plot of a portion of the polymeric structure drawn at 30% probability displacement ellipsoids. Symmetry codes: i: $-x + 1, y - 1/2, -z + 3/2$; ii: $-x, y - 1/2, -z + 3/2$.

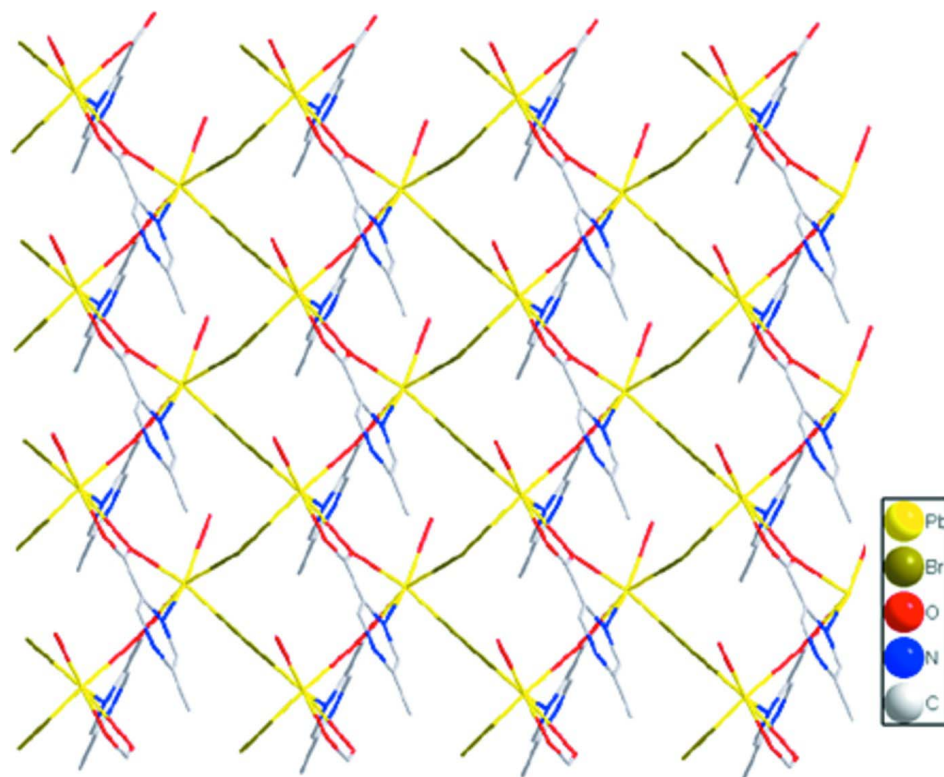


Figure 2
Layer structures.

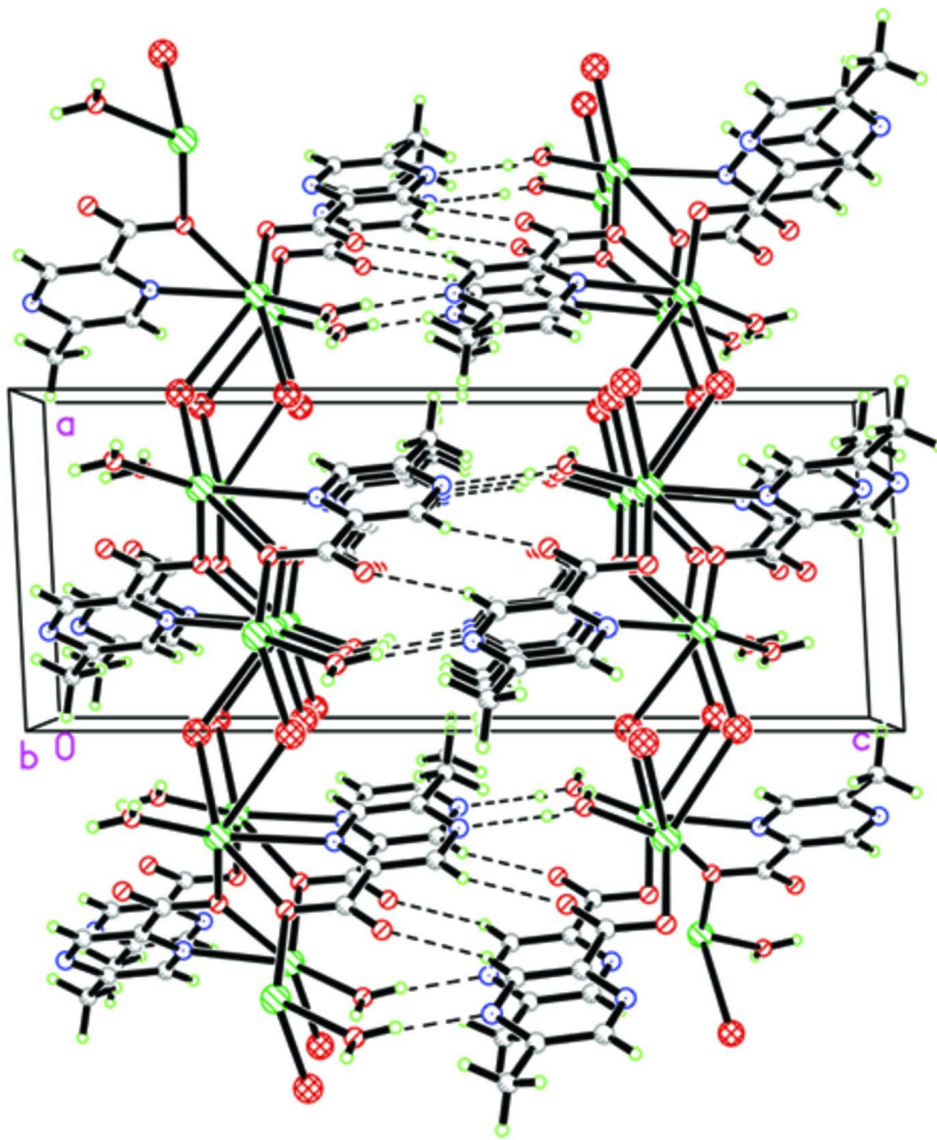


Figure 3

Three-dimensional supramolecular structure. Dashed lines represent O—H \cdots N and weak O—H \cdots Br hydrogen bonds.

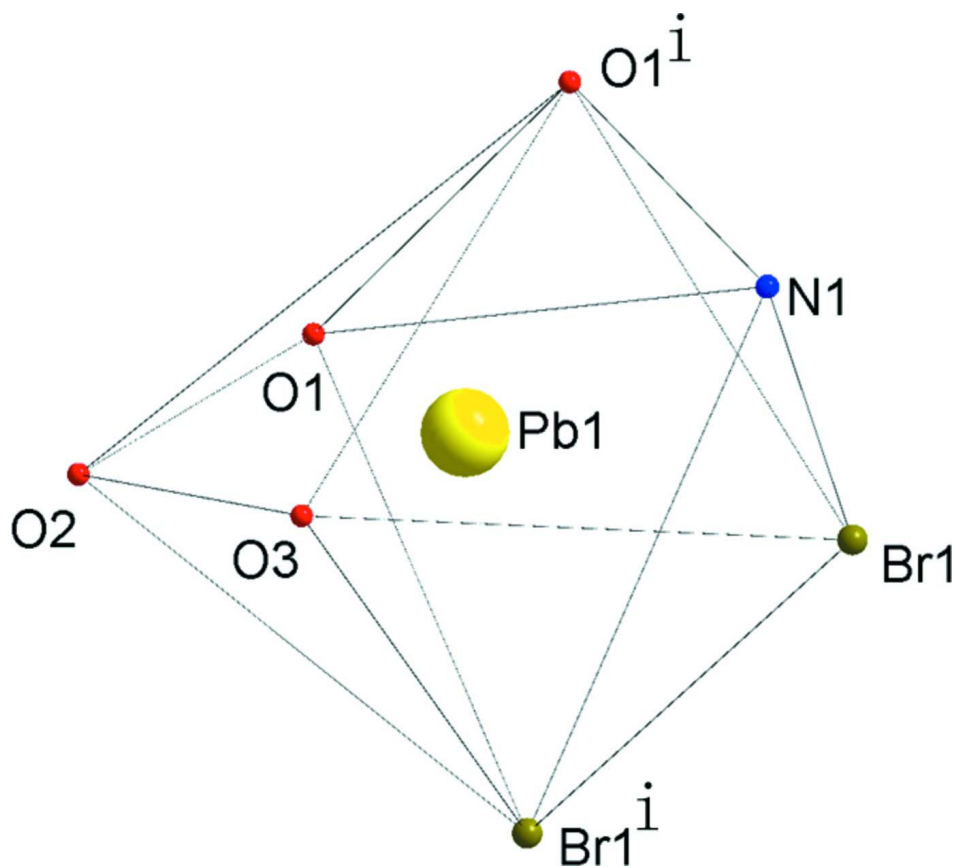


Figure 4

Ψ -Dodecahedral geometry of lead(II).

Poly[aqua- μ -bromido-(μ_2 -5-methylpyrazine-2-carboxylato- $\kappa^4 N^1, O^2:O^2, O^2$)lead(II)]

Crystal data

[PbBr(C₆H₅N₂O₂)(H₂O)]

$M_r = 442.24$

Monoclinic, $P2_1/c$

Hall symbol: -P 2ybc

$a = 7.5493$ (10) Å

$b = 6.6775$ (9) Å

$c = 19.335$ (3) Å

$\beta = 92.884$ (2)°

$V = 973.5$ (2) Å³

$Z = 4$

$F(000) = 792$

$D_x = 3.018$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 3510 reflections

$\theta = 3.3$ – 28.2 °

$\mu = 21.41$ mm⁻¹

$T = 296$ K

Block, colourless

$0.15 \times 0.14 \times 0.13$ mm

Data collection

Bruker SMART APEX CCD area-detector
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

phi and ω scans

Absorption correction: multi-scan

(*SADABS*; Sheldrick, 1996)

$T_{\min} = 0.142$, $T_{\max} = 0.167$

5123 measured reflections

1904 independent reflections

1747 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.039$

$\theta_{\max} = 26.0$ °, $\theta_{\min} = 3.2$ °

$h = -9 \rightarrow 5$

$k = -8 \rightarrow 8$

$l = -23 \rightarrow 22$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.033$
 $wR(F^2) = 0.091$
 $S = 1.05$
 1904 reflections
 120 parameters

0 restraints
 H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0596P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.032$
 $\Delta\rho_{\max} = 2.26 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -2.20 \text{ e } \text{\AA}^{-3}$

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R -factors(gt) etc. and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
Pb1	0.29327 (3)	-0.20899 (3)	0.782606 (12)	0.02328 (17)
Br1	0.00888 (10)	0.08548 (11)	0.80760 (4)	0.0386 (2)
O1	0.5113 (7)	0.0867 (8)	0.7933 (3)	0.0351 (12)
O2	0.4664 (8)	-0.0254 (8)	0.8984 (3)	0.0427 (13)
O3	0.2211 (10)	-0.5006 (9)	0.8682 (3)	0.067 (2)
H3A	0.2481	-0.4963	0.9114	0.100*
H3B	0.1828	-0.6189	0.8608	0.100*
N1	0.6818 (7)	0.4165 (8)	0.8455 (3)	0.0274 (12)
N2	0.7181 (8)	0.4728 (9)	0.9872 (3)	0.0322 (13)
C1	0.6189 (9)	0.2822 (9)	0.8890 (4)	0.0245 (15)
C2	0.6373 (10)	0.3105 (10)	0.9598 (4)	0.0302 (16)
H2	0.5925	0.2146	0.9891	0.036*
C3	0.7783 (9)	0.6113 (9)	0.9431 (4)	0.0298 (15)
C4	0.7588 (9)	0.5805 (9)	0.8721 (4)	0.0285 (15)
H4	0.8006	0.6771	0.8423	0.034*
C5	0.5250 (9)	0.0994 (10)	0.8587 (4)	0.0280 (15)
C6	0.8627 (14)	0.7958 (11)	0.9749 (6)	0.049 (2)
H6A	0.7859	0.8517	1.0079	0.073*
H6B	0.8818	0.8924	0.9392	0.073*
H6C	0.9743	0.7609	0.9977	0.073*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Pb1	0.0230 (2)	0.0234 (2)	0.0233 (2)	-0.00092 (7)	0.00028 (12)	0.00041 (8)
Br1	0.0362 (4)	0.0338 (4)	0.0451 (5)	0.0095 (3)	-0.0054 (3)	0.0001 (3)
O1	0.036 (3)	0.042 (3)	0.027 (3)	-0.017 (2)	-0.005 (2)	-0.004 (2)
O2	0.056 (4)	0.038 (3)	0.034 (3)	-0.015 (2)	-0.002 (2)	0.009 (2)

O3	0.119 (7)	0.049 (4)	0.031 (3)	-0.039 (4)	-0.019 (3)	0.008 (3)
N1	0.028 (3)	0.026 (3)	0.028 (3)	-0.001 (2)	-0.001 (2)	-0.002 (2)
N2	0.040 (4)	0.030 (3)	0.026 (3)	-0.004 (2)	-0.001 (2)	-0.005 (2)
C1	0.021 (3)	0.026 (4)	0.027 (4)	0.000 (2)	-0.002 (3)	0.000 (3)
C2	0.031 (4)	0.031 (4)	0.029 (4)	-0.001 (3)	0.003 (3)	0.001 (3)
C3	0.030 (4)	0.024 (3)	0.035 (4)	0.000 (3)	-0.003 (3)	-0.003 (3)
C4	0.031 (4)	0.023 (3)	0.031 (4)	-0.002 (3)	0.001 (3)	-0.001 (3)
C5	0.025 (3)	0.030 (3)	0.029 (4)	-0.006 (3)	0.003 (3)	-0.003 (3)
C6	0.059 (6)	0.033 (5)	0.055 (6)	-0.011 (3)	0.004 (5)	-0.008 (4)

Geometric parameters (Å, °)

Pb1—O1 ⁱ	2.531 (5)	N1—C1	1.334 (8)
Pb1—O1	2.572 (5)	N1—Pb1 ^{iv}	2.631 (5)
Pb1—N1 ⁱ	2.631 (6)	N2—C2	1.340 (9)
Pb1—O3	2.631 (6)	N2—C3	1.353 (9)
Pb1—Br1	2.9688 (8)	C1—C2	1.381 (11)
Pb1—Br1 ⁱⁱ	3.1190 (9)	C1—C5	1.514 (9)
Br1—Pb1 ⁱⁱⁱ	3.1190 (9)	C2—H2	0.9300
O1—C5	1.268 (8)	C3—C4	1.388 (10)
O1—Pb1 ^{iv}	2.531 (5)	C3—C6	1.504 (10)
O2—C5	1.230 (8)	C4—H4	0.9300
O3—H3A	0.8501	C6—H6A	0.9600
O3—H3B	0.8510	C6—H6B	0.9600
N1—C4	1.331 (9)	C6—H6C	0.9600
O1 ⁱ —Pb1—O1	94.06 (8)	C1—N1—Pb1 ^{iv}	115.1 (4)
O1 ⁱ —Pb1—N1 ⁱ	63.56 (16)	C2—N2—C3	117.6 (6)
O1—Pb1—N1 ⁱ	75.81 (17)	N1—C1—C2	120.8 (6)
O1 ⁱ —Pb1—O3	96.4 (2)	N1—C1—C5	118.2 (6)
O1—Pb1—O3	132.06 (18)	C2—C1—C5	121.0 (6)
N1 ⁱ —Pb1—O3	148.79 (18)	N2—C2—C1	121.6 (6)
O1 ⁱ —Pb1—Br1	153.98 (12)	N2—C2—H2	119.2
O1—Pb1—Br1	86.76 (12)	C1—C2—H2	119.2
N1 ⁱ —Pb1—Br1	91.65 (12)	N2—C3—C4	120.0 (6)
O3—Pb1—Br1	102.33 (18)	N2—C3—C6	116.8 (7)
O1 ⁱ —Pb1—Br1 ⁱⁱ	82.54 (13)	C4—C3—C6	123.2 (7)
O1—Pb1—Br1 ⁱⁱ	146.05 (12)	N1—C4—C3	121.8 (6)
N1 ⁱ —Pb1—Br1 ⁱⁱ	72.46 (13)	N1—C4—H4	119.1
O3—Pb1—Br1 ⁱⁱ	81.80 (15)	C3—C4—H4	119.1
Br1—Pb1—Br1 ⁱⁱ	82.410 (17)	O2—C5—O1	124.3 (6)
Pb1—Br1—Pb1 ⁱⁱⁱ	135.64 (3)	O2—C5—C1	118.7 (6)
C5—O1—Pb1 ^{iv}	121.4 (4)	O1—C5—C1	117.0 (5)
C5—O1—Pb1	98.7 (4)	C3—C6—H6A	109.5
Pb1 ^{iv} —O1—Pb1	139.2 (2)	C3—C6—H6B	109.5
Pb1—O3—H3A	123.1	H6A—C6—H6B	109.5
Pb1—O3—H3B	131.3	C3—C6—H6C	109.5
H3A—O3—H3B	105.1	H6A—C6—H6C	109.5

C4—N1—C1	118.2 (6)	H6B—C6—H6C	109.5
C4—N1—Pb1 ^{iv}	125.1 (4)		
O1 ⁱ —Pb1—Br1—Pb1 ⁱⁱⁱ	13.5 (3)	Pb1 ^{iv} —N1—C1—C5	-15.9 (7)
O1—Pb1—Br1—Pb1 ⁱⁱⁱ	106.16 (12)	C3—N2—C2—C1	1.4 (10)
N1 ⁱ —Pb1—Br1—Pb1 ⁱⁱⁱ	30.47 (13)	N1—C1—C2—N2	0.1 (11)
O3—Pb1—Br1—Pb1 ⁱⁱⁱ	-121.43 (15)	C5—C1—C2—N2	-179.1 (6)
Br1 ⁱⁱ —Pb1—Br1—Pb1 ⁱⁱⁱ	-41.59 (4)	C2—N2—C3—C4	-1.4 (10)
O1 ⁱ —Pb1—O1—C5	-126.3 (4)	C2—N2—C3—C6	177.8 (7)
N1 ⁱ —Pb1—O1—C5	172.3 (4)	C1—N1—C4—C3	1.6 (10)
O3—Pb1—O1—C5	-23.9 (5)	Pb1 ^{iv} —N1—C4—C3	-163.3 (5)
Br1—Pb1—O1—C5	79.8 (4)	N2—C3—C4—N1	-0.1 (10)
Br1 ⁱⁱ —Pb1—O1—C5	151.1 (3)	C6—C3—C4—N1	-179.3 (7)
O1 ⁱ —Pb1—O1—Pb1 ^{iv}	43.5 (3)	Pb1 ^{iv} —O1—C5—O2	-162.2 (6)
N1 ⁱ —Pb1—O1—Pb1 ^{iv}	-17.9 (3)	Pb1—O1—C5—O2	10.0 (8)
O3—Pb1—O1—Pb1 ^{iv}	145.9 (4)	Pb1 ^{iv} —O1—C5—C1	18.3 (8)
Br1—Pb1—O1—Pb1 ^{iv}	-110.4 (3)	Pb1—O1—C5—C1	-169.5 (5)
Br1 ⁱⁱ —Pb1—O1—Pb1 ^{iv}	-39.1 (5)	N1—C1—C5—O2	-179.9 (6)
C4—N1—C1—C2	-1.5 (9)	C2—C1—C5—O2	-0.7 (10)
Pb1 ^{iv} —N1—C1—C2	164.8 (5)	N1—C1—C5—O1	-0.4 (9)
C4—N1—C1—C5	177.7 (6)	C2—C1—C5—O1	178.8 (6)

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

Hydrogen-bond geometry (Å, °)

<i>D—H...A</i>	<i>D—H</i>	<i>H...A</i>	<i>D...A</i>	<i>D—H...A</i>
O3—H3A...N2 ^v	0.85	1.97	2.816 (9)	173
O3—H3B...Br1 ^{vi}	0.85	2.56	3.378 (6)	161

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