

Redetermination of *catena*-poly[[chloridolead(II)]- μ_2 -chlorido-di- μ_2 -thiourea- κ^4 S:S] at 100 K

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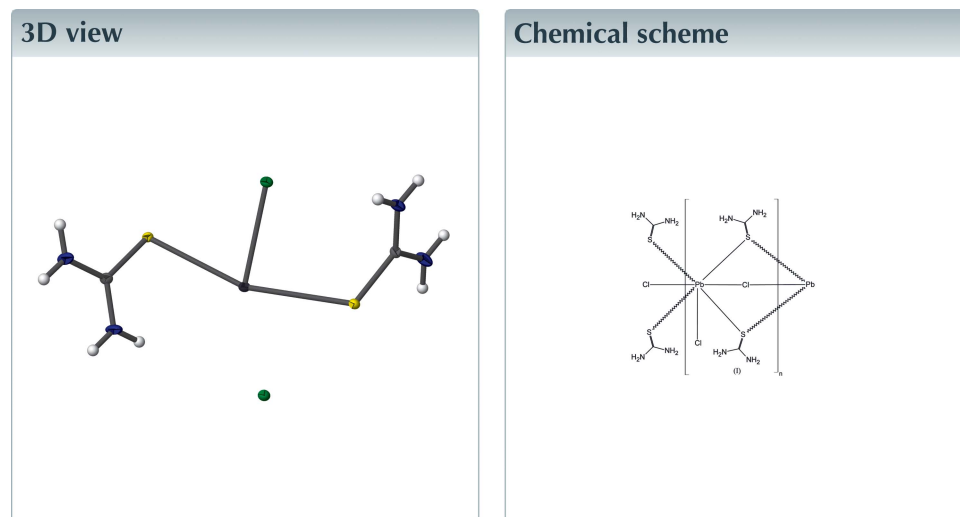
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Keywords: crystal structure; redetermination; polymeric lead thiourea salt.

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Structural data: full structural data are available from iucrdata.iucr.org

Although the structure refinement (CCD data at 100 K) of the polymeric lead(II) thiourea complex, $[\text{PbCl}_2 \cdot 2\text{tu}]_n$ where ($\text{tu} = \text{SCN}_2\text{H}_4$), basically confirmed the previous study based on integrated Weissenberg data recorded at room temperature [Nardelli & Fava (1959). *Acta Cryst.* **12**, 727–732]; all atomic positions could be determined with significantly higher precision and accuracy. In addition, all H atoms could be located from difference maps, revealing details of the hydrogen-bonding scheme.



Structure description

A survey in the Cambridge Structural Database (Groom *et al.*, 2016) shows that the interaction of thiourea (tu) with lead(II) can result in a variety of different compounds and types of coordination. A sixfold coordination of lead by thiourea is found in triclinic $\text{Pb}(\text{ClO}_4)_2 \cdot 6\text{tu}$ (Goldberg & Herbstein, 1972), sevenfold coordination in $\text{PbCl}_2 \cdot 2\text{tu}$ (Nardelli & Fava, 1959) – which is redetermined in the present work – and eightfold coordination in $\text{PbH}(\text{COO})_2 \cdot 4\text{tu} \cdot \text{H}_2\text{O}$ (Goldberg & Herbstein, 1973) or $\text{Pb}(\text{CHOO})_2 \cdot 2\text{tu}$ (Nardelli *et al.*, 1960).

The crystal structure of the title compound comprises an infinite polymeric chain propagating along the b -axis direction (Fig. 1) in which the Pb^{II} ions are linked by six bridging atoms, namely four S atoms from two pairs of symmetry-related thiourea ligands and by two symmetry-related Cl^- anions. The distorted sevenfold coordination is completed by another terminal Cl^- anion. The resulting $[\text{PbS}_4\text{Cl}_3]$ polyhedron can be derived from a distorted trigonal prism formed by four S and two Cl atoms that is capped on one of the lateral faces by another Cl atom. In comparison with the previous determination of the title compound (Nardelli & Fava, 1959) that was based on integrated

Table 1

Comparison of bond lengths (Å) in the current and the previous (Nardelli & Fava, 1959) refinement of poly[μ_2 -chlorido-chloridobis(μ_2 -thiourea- κ^2 S:S)lead(II)].

For the previous refinement: $a = 21.20$ (4), $b = 4.06$ (1), $c = 12.02$ (2) Å; $R = 0.114$.

Bond	Current refinement	Previous refinement
Pb—Cl1	2.7451 (8)	2.75 (4)
Pb—Cl2	3.1708 (7)	3.17 (3)
Pb—Cl1 ⁱ	3.1635 (7)	3.28 (3)
Pb—S1	3.0057 (7)	3.02 (3)
Pb—S1 ⁱⁱⁱ	3.0494 (7)	3.04 (3)
Pb—S2 ⁱ	2.9923 (7)	3.10 (3)
Pb—S2	2.9494 (7)	2.92 (3)
S1—C1	1.732 (3)	1.68 (9)
S2—C2	1.739 (3)	1.78 (17)
C1—N1	1.322 (4)	1.40 (12)
C1—N2	1.328 (4)	1.35 (14)
C2—N3	1.320 (4)	1.32 (20)
C2—N4	1.323 (4)	1.34 (17)

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

Table 2

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N4—H1A \cdots Cl1 ⁱ	0.86	2.53	3.299 (3)	150
N4—H1B \cdots Cl2 ⁱⁱ	0.86	2.50	3.336 (3)	163
N3—H2A \cdots Cl1 ⁱ	0.86	2.78	3.494 (4)	142
N3—H2A \cdots S1 ⁱⁱⁱ	0.86	2.85	3.421 (3)	126
N3—H2B \cdots Cl1 ^{iv}	0.86	2.44	3.285 (3)	168
N1—H3A \cdots Cl2 ^v	0.86	2.72	3.480 (4)	149
N1—H3A \cdots S2 ^v	0.86	2.75	3.414 (3)	135
N1—H3B \cdots Cl2 ^{vi}	0.86	2.37	3.232 (4)	175
N2—H4A \cdots Cl2 ^v	0.86	2.49	3.304 (3)	159
N2—H4B \cdots Cl1 ^{vii}	0.86	2.44	3.275 (3)	164

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

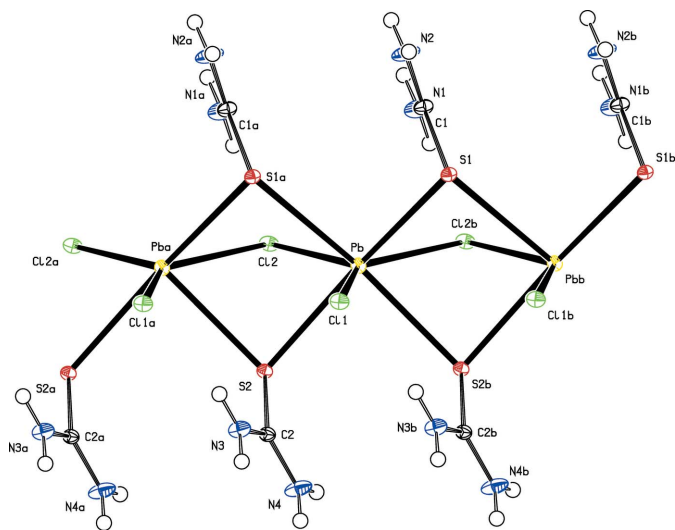


Figure 1

The polymeric chain in the crystal structure of the title compound. Displacement ellipsoids are drawn at the 30% probability level. H atoms are represented as small spheres of arbitrary radii. [Symmetry codes: (a) $x, y + 1, z$; (b) $x, y - 1, z$.]

Table 3

Experimental details.

Crystal data	[PbCl ₂ (CH ₄ N ₂ S) ₂]
Chemical formula	430.33
M_r	Orthorhombic, $Pna2_1$
Crystal system, space group	100
Temperature (K)	21.1951 (6), 4.0280 (1), 11.9433 (3)
a, b, c (Å)	1019.65 (5)
V (Å ³)	4
Z	Mo $K\alpha$
Radiation type	17.43
μ (mm ⁻¹)	0.46 × 0.17 × 0.14
Crystal size (mm)	
Data collection	Bruker APEXII CCD detector
Diffractometer	Multi-scan (SADABS; Krause <i>et al.</i> , 2015)
Absorption correction	0.040, 0.087
T_{\min}, T_{\max}	23237, 6206, 5867
No. of measured, independent and observed [$I > 2\sigma(I)$] reflections	0.032
R_{int}	0.904
$(\sin \theta/\lambda)_{\text{max}}$ (Å ⁻¹)	
Refinement	
$R[F^2 > 2\sigma(F^2)], wR(F^2), S$	0.018, 0.041, 1.04
No. of reflections	6206
No. of parameters	101
No. of restraints	1
H-atom treatment	H-atom parameters constrained
$\Delta\rho_{\text{max}}, \Delta\rho_{\text{min}}$ (e Å ⁻³)	1.18, -2.38
Absolute structure	Refined as an inversion twin
Absolute structure parameter	0.315 (4)

Computer programs: APEX2 and SAINT (Bruker, 2009), SHELXT (Sheldrick, 2015a), SHELXL2014 (Sheldrick, 2015b), PLATON (Spek, 2009) and publCIF (Westrip, 2010).

Weissenberg data (room-temperature measurement), the current redetermination reveals not only a higher precision but also a significantly higher accuracy, in particular for the C=S, C—C and C—N bond lengths (Table 1). In contrast to the previous study, the H atoms could be determined in the

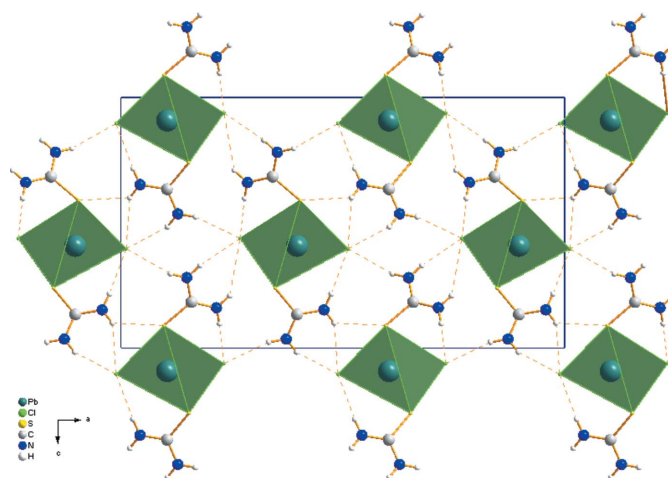


Figure 2

The crystal packing of the title compound viewed along the b axis. The coordination sphere around the Pb^{II} atom is given in the polyhedral representation; N—H \cdots Cl and N—H \cdots S hydrogen bonds are shown as dashed lines (see Table 2 for details).

current study. All of them are involved in hydrogen-bonding interactions. Intra-chain N—H···Cl hydrogen bonds as well as interchain N—H···Cl and N—H···S hydrogen bonds (Table 2) are observed in the crystal, leading to a three-dimensional network structure (Fig. 2).

Synthesis and crystallization

To obtain the title compound, (diaminomethylidene)-sulfonium chloride–thiourea (3/2) (Zouihri, 2012) (1 mmol) in ethanol (10 ml) was added dropwise to an aqueous solution (5 ml) of lead chlorate (2 mmol). The resulting solution was allowed to stand at room temperature. After two weeks, colourless crystals with good quality were obtained from the filtrate and dried in air.

Refinement

Crystal data, data collection and structure refinement details are summarized in Table 3. The same atom labelling as in the previous study (Nardelli & Fava, 1959) was used for better

comparison (Table 1). All hydrogen atoms could be localized in difference Fourier syntheses. The structure was refined as an inversion twin (Table 3), using 5385 Friedel pairs. Reflections (200) and (201) were omitted from the refinement due to obstruction from the beam stop.

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full crystallographic data

IUCrData (2016). 1, x161745 [https://doi.org/10.1107/S2414314616017454]

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catena-Poly[[chloridolead(II)]- μ_2 -chlorido-di- μ_2 -thiourea- κ^4 S:S]

Crystal data

[PbCl₂(CH₄N₂S)₂]

$M_r = 430.33$

Orthorhombic, *Pna*2₁

$a = 21.1951$ (6) Å

$b = 4.0280$ (1) Å

$c = 11.9433$ (3) Å

$V = 1019.65$ (5) Å³

$Z = 4$

$F(000) = 784$

$D_x = 2.803$ Mg m⁻³

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

Cell parameters from 318 reflections

$\theta = 1.5$ – 28.3°

$\mu = 17.43$ mm⁻¹

$T = 100$ K

Prism, colourless

$0.46 \times 0.17 \times 0.14$ mm

Data collection

Bruker APEXII CCD detector
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

ω and φ scans

Absorption correction: multi-scan
(*SADABS*; Krause *et al.*, 2015)

$T_{\min} = 0.040$, $T_{\max} = 0.087$

23237 measured reflections

6206 independent reflections

5867 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.032$

$\theta_{\max} = 40.0^\circ$, $\theta_{\min} = 2.6^\circ$

$h = -38 \rightarrow 36$

$k = -7 \rightarrow 5$

$l = -21 \rightarrow 21$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.018$

$wR(F^2) = 0.041$

$S = 1.04$

6206 reflections

101 parameters

1 restraint

Hydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0121P)^2]$

where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} = 0.001$

$\Delta\rho_{\max} = 1.18$ e Å⁻³

$\Delta\rho_{\min} = -2.38$ e Å⁻³

Absolute structure: Refined as an inversion twin

Absolute structure parameter: 0.315 (4)

Special details

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

Refinement. Refined as a 2-component inversion twin

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

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
Pb	0.10368 (2)	0.29058 (2)	0.09284 (2)	0.00922 (2)
Cl1	0.23177 (4)	0.24529 (17)	0.06240 (7)	0.01151 (10)
Cl2	-0.01146 (3)	-0.21081 (15)	0.10426 (7)	0.01147 (13)
S2	0.09634 (4)	-0.20155 (16)	-0.08955 (7)	0.00909 (11)
S1	0.15296 (4)	0.78245 (15)	0.26068 (7)	0.00912 (11)
C1	0.10345 (13)	0.6395 (9)	0.3655 (3)	0.0122 (4)
C2	0.15858 (15)	-0.1690 (6)	-0.1838 (2)	0.0102 (4)
N3	0.21531 (14)	-0.2855 (6)	-0.1607 (3)	0.0145 (4)
H2A	0.2453	-0.2656	-0.2088	0.017*
H2B	0.2223	-0.3812	-0.0975	0.017*
N1	0.04268 (13)	0.5872 (8)	0.3479 (3)	0.0187 (5)
H3A	0.0195	0.5066	0.4004	0.022*
H3B	0.0263	0.6338	0.2839	0.022*
N2	0.12793 (14)	0.5659 (8)	0.4648 (2)	0.0185 (5)
H4A	0.1043	0.4854	0.5166	0.022*
H4B	0.1674	0.5988	0.4772	0.022*
N4	0.14795 (14)	-0.0218 (8)	-0.2809 (2)	0.0196 (5)
H1A	0.1780	-0.0023	-0.3289	0.024*
H1B	0.1110	0.0543	-0.2961	0.024*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Pb	0.00719 (4)	0.01007 (3)	0.01040 (4)	0.00009 (2)	0.00000 (5)	-0.00047 (9)
Cl1	0.0079 (3)	0.0156 (2)	0.0110 (2)	0.00059 (18)	0.0001 (2)	-0.00040 (17)
Cl2	0.0080 (2)	0.0163 (2)	0.0101 (4)	-0.00025 (15)	0.0003 (2)	-0.0017 (2)
S2	0.0074 (3)	0.0108 (2)	0.0090 (3)	-0.00018 (17)	0.0008 (2)	-0.00054 (19)
S1	0.0080 (3)	0.0107 (2)	0.0086 (3)	-0.00211 (18)	0.0000 (2)	0.0005 (2)
C1	0.0103 (12)	0.0157 (10)	0.0107 (11)	-0.0031 (8)	0.0007 (8)	0.0012 (9)
C2	0.0097 (11)	0.0118 (9)	0.0091 (10)	0.0014 (7)	0.0004 (9)	-0.0004 (7)
N3	0.0087 (11)	0.0224 (11)	0.0125 (11)	0.0045 (8)	0.0014 (9)	0.0029 (8)
N1	0.0075 (10)	0.0341 (14)	0.0146 (11)	-0.0044 (9)	-0.0003 (9)	0.0046 (11)
N2	0.0114 (12)	0.0343 (15)	0.0099 (10)	-0.0059 (10)	-0.0019 (9)	0.0063 (9)
N4	0.0127 (12)	0.0342 (14)	0.0119 (11)	0.0076 (10)	0.0033 (9)	0.0081 (10)

Geometric parameters (\AA , $^\circ$)

Pb—Cl1	2.7451 (8)	C1—N2	1.328 (4)
Pb—S2	2.9494 (7)	C2—N3	1.320 (4)
Pb—S2 ⁱ	2.9923 (7)	C2—N4	1.323 (4)
Pb—S1	3.0057 (7)	N3—H2A	0.8600
Pb—S1 ⁱⁱ	3.0494 (7)	N3—H2B	0.8600
Pb—Cl2	3.1708 (7)	N1—H3A	0.8600
S2—C2	1.739 (3)	N1—H3B	0.8600
S2—Pb ⁱⁱ	2.9924 (7)	N2—H4A	0.8600

S1—C1	1.732 (3)	N2—H4B	0.8600
S1—Pb ⁱ	3.0493 (7)	N4—H1A	0.8600
C1—N1	1.322 (4)	N4—H1B	0.8600
C11—Pb—S2	84.82 (2)	Pb—S1—Pb ⁱ	83.40 (2)
C11—Pb—S2 ⁱ	90.03 (2)	N1—C1—N2	119.1 (3)
S2—Pb—S2 ⁱ	85.36 (2)	N1—C1—S1	121.9 (3)
C11—Pb—S1	77.79 (2)	N2—C1—S1	118.9 (2)
S2—Pb—S1	162.54 (2)	N3—C2—N4	119.9 (3)
S2 ⁱ —Pb—S1	93.035 (19)	N3—C2—S2	122.0 (2)
C11—Pb—S1 ⁱⁱ	72.76 (2)	N4—C2—S2	118.2 (2)
S2—Pb—S1 ⁱⁱ	93.01 (2)	C2—N3—H2A	120.0
S2 ⁱ —Pb—S1 ⁱⁱ	162.79 (2)	C2—N3—H2B	120.0
S1—Pb—S1 ⁱⁱ	83.40 (2)	H2A—N3—H2B	120.0
C11—Pb—Cl2	136.433 (18)	C1—N1—H3A	120.0
S2—Pb—Cl2	64.09 (2)	C1—N1—H3B	120.0
S2 ⁱ —Pb—Cl2	115.26 (2)	H3A—N1—H3B	120.0
S1—Pb—Cl2	131.19 (2)	C1—N2—H4A	120.0
S1 ⁱⁱ —Pb—Cl2	78.92 (2)	C1—N2—H4B	120.0
C2—S2—Pb	112.77 (10)	H4A—N2—H4B	120.0
C2—S2—Pb ⁱⁱ	118.89 (10)	C2—N4—H1A	120.0
Pb—S2—Pb ⁱⁱ	85.36 (2)	C2—N4—H1B	120.0
C1—S1—Pb	92.99 (11)	H1A—N4—H1B	120.0
C1—S1—Pb ⁱ	119.38 (11)		

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

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N4—H1A \cdots C11 ⁱⁱⁱ	0.86	2.53	3.299 (3)	150
N4—H1B \cdots C12 ^{iv}	0.86	2.50	3.336 (3)	163
N3—H2A \cdots C11 ⁱⁱⁱ	0.86	2.78	3.494 (4)	142
N3—H2A \cdots S1 ^v	0.86	2.85	3.421 (3)	126
N3—H2B \cdots C11 ⁱⁱ	0.86	2.44	3.285 (3)	168
N1—H3A \cdots C12 ^{vi}	0.86	2.72	3.480 (4)	149
N1—H3A \cdots S2 ^{vi}	0.86	2.75	3.414 (3)	135
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