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Dieter Schollmeyer^{b*}^aInstitut für Anorganische Chemie und Analytische Chemie der Universität Mainz, Duesbergweg 10-14, 55099 Mainz, Germany, and ^bInstitut für Organische Chemie der Universität Mainz, Duesbergweg 10-14, 55099 Mainz, Federal Republic of Germany

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Key indicators

Single-crystal X-ray study
 $T = 298\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.015\text{ \AA}$
 R factor = 0.051
 wR factor = 0.149
Data-to-parameter ratio = 15.6For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.Poly[bis(μ -pentafluorobenzenethiolato)lead(II)]

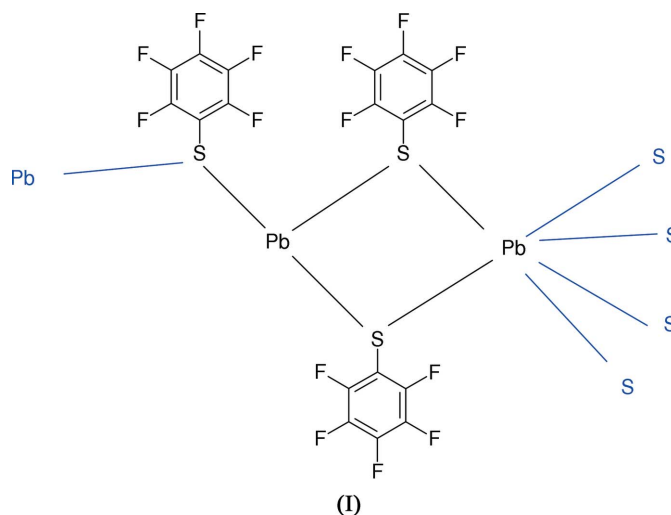
The title compound, $[\text{Pb}(\text{SC}_6\text{F}_5)_2]_n$, exhibits a layered structure containing two differently hexacoordinated Pb^{II} atoms. One is situated on a centre of inversion and is coordinated by six S atoms, its valence lone pair of electrons being consequently stereochemically inactive. The other Pb atom is pyramidally coordinated by three S atoms and forms three contacts to adjacent F atoms, the stereochemically active lone pair pointing towards the F_3 plane.

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Comment

Pb^{II} thiolates without additional donor atoms tend to form thiolate-bridged polymeric solid state structures and hence are soluble only in strongly polar solvents (Dance, 1986). The title compound, (I), which is of interest for synthetic purposes (Peach & Spinney, 1971; Redón *et al.*, 2001; Morales-Morales *et al.*, 2001), is very soluble in acetone and methanol. It was thus of interest to explore its crystal structure and to see whether intra- or intermolecular $\text{Pb}\cdots\text{F}$ contacts are present.



Two quite different hexacoordinate Pb^{II} centres are present in (I). Pb1 is situated on an inversion centre, its valence lone pair being consequently stereochemically inactive (Fig. 1). It is coordinated by six S atoms and exhibits a distorted octahedral coordination geometry, with $\text{Pb1}-\text{S}$ distances between 2.971 (2) and 3.036 (2) Å and *cis* $\text{S}-\text{Pb1}-\text{S}$ angles between 66.49 (6) and 113.51 (6)°.

Atom Pb2 is coordinated by three SC_6F_5 ligands *via* their S atoms. Each ligand additionally forms a $\text{Pb}\cdots\text{F}$ contact with the F atom in the position *ortho* to the S atom (Fig. 2). The $\text{Pb2}-\text{S}$ distances range between 2.704 (2) and 2.737 (2) Å and are thus significantly shorter than the $\text{Pb1}-\text{S}$ bonds, while the

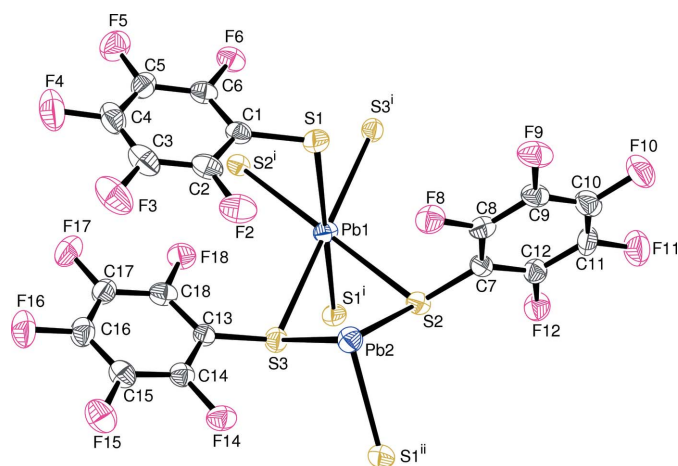


Figure 1
Detail of (I), showing the Pb coordination environments (30% displacement ellipsoids). [Symmetry codes: (i) $1 - x, 1 - y, 1 - z$; (ii) $1 - x, y - \frac{1}{2}, \frac{1}{2} - z$.]

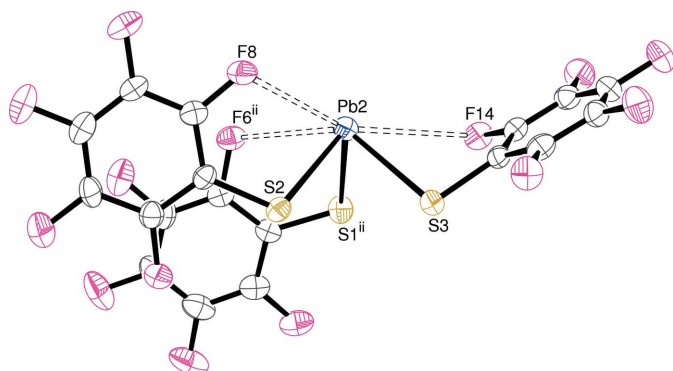


Figure 2
Environment of atom Pb2 in (I), showing the Pb...F interactions as dashed lines (30% displacement ellipsoids). The symmetry codes are as in Fig. 1.

Pb...F contacts are rather long [2.968 (6)–3.245 (6) Å]. The PbS₃ unit exhibits a trigonal-pyramidal configuration with a fairly narrow spread of S–Pb2–S angles between 75.61 (7) and 86.77 (8)°. The stereochemically active lone pair of electrons points towards the F₃ plane, from which Pb2 is displaced by 0.339 (3) Å.

Atoms Pb1 and Pb2 are connected via μ^2 -S bridges and form a sheet structure consisting of PbS₆ octahedra and PbS₃ pyramids sharing corners and edges (Fig. 3). It is interesting to compare the crystal structure of (I) to that of Pb(SAr)₂ (Ar = C₆H₂’Bu₃-2,4,6; Hitchcock *et al.* 1983). In both, pyramidal PbS₃ units form μ^2 -S bridges to another crystallographically different Pb atom. In Pb(SAr)₂, this second Pb atom exhibits a pseudo-trigonal-bipyramidal PbS₄ coordination mode, with a stereochemically active lone pair in an equatorial position, and is thus different from the octahedrally coordinated Pb1 in Pb(SC₆F₅)₂.

The Pb–S bonds in (I) are longer [mean Pb–S = 3.012 (3) Å for the six-coordinate site and mean Pb–S = 2.721 (3) Å for the three-coordinate site] and hence weaker

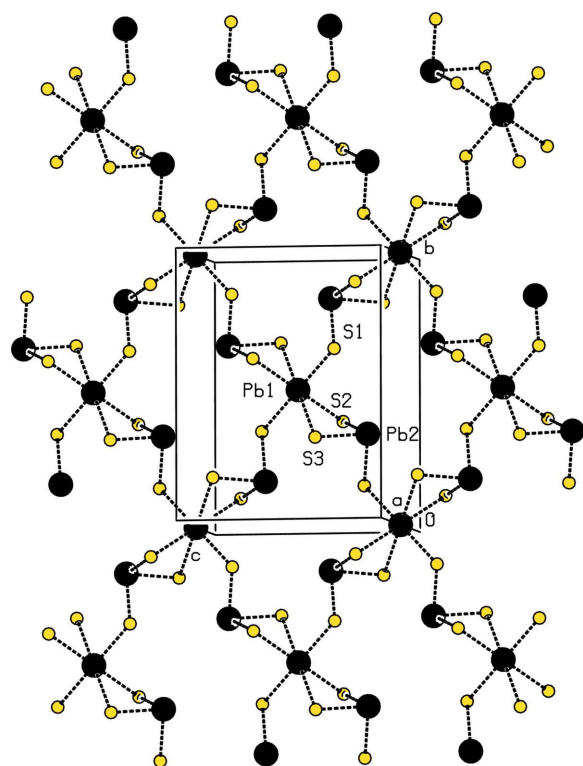


Figure 3
Packing diagram of the Pb and S atoms in (I), showing the sheet structure.

than those in Pb(SAr)₂ [mean Pb–S = 2.788 (3) Å for the four-coordinate site and mean Pb–S = 2.669 (3) Å for the three-coordinate site]. This may explain the enhanced solubility of Pb(SC₆F₅)₂ in acetone and methanol, despite the polymeric nature of its structure in the solid state.

Experimental

Compound (I) was prepared according to the literature procedure of Peach (1968). Its purity was confirmed by elemental (C and S) analysis and ¹⁹F NMR spectroscopy. Crystals suitable for single-crystal X-ray diffraction were obtained by cooling a saturated solution of the compound in acetone from room temperature to 278 K.

Crystal data

[Pb(C ₆ F ₅ S) ₂]	Z = 6
<i>M_r</i> = 605.43	<i>D_x</i> = 2.824 Mg m ^{−3}
Monoclinic, <i>P</i> 2 ₁ / <i>c</i>	Mo <i>K</i> α radiation
<i>a</i> = 14.7217 (5) Å	μ = 12.25 mm ^{−1}
<i>b</i> = 13.7581 (5) Å	<i>T</i> = 298 (2) K
<i>c</i> = 10.5960 (4) Å	Plate, yellow
β = 95.647 (1)°	0.28 × 0.24 × 0.09 mm
<i>V</i> = 2135.73 (13) Å ³	

Data collection

Bruker SMART CCD diffractometer	32252 measured reflections
ω scans	5306 independent reflections
Absorption correction: multi-scan (SORTAV; Blessing, 1995)	3532 reflections with <i>I</i> > 2 σ (<i>I</i>)
<i>T</i> _{min} = 0.03, <i>T</i> _{max} = 0.30	<i>R</i> _{int} = 0.161
(expected range = 0.033–0.332)	θ _{max} = 28.3°

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.051$ $wR(F^2) = 0.149$ $S = 1.01$

5306 reflections

340 parameters

$$w = 1/[\sigma^2(F_o^2) + (0.076P)^2 + 4.1862P]$$

$$\text{where } P = (F_o^2 + 2F_c^2)/3$$

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

$$\Delta\rho_{\max} = 2.22 \text{ e } \text{\AA}^{-3}$$

$$\Delta\rho_{\min} = -1.58 \text{ e } \text{\AA}^{-3}$$

Table 1Selected bond lengths (\AA).

Pb1—S1	2.971 (2)	Pb2—S2	2.704 (2)
Pb1—S2	3.030 (2)	Pb2—S3	2.721 (2)
Pb1—S3	3.036 (2)	Pb2—S1 ⁱ	2.737 (3)

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

The highest peak is located 0.87 \AA from atom Pb2 and the deepest hole is located 0.69 \AA from the same atom. The high value of R_{int} is due to the poor quality of the crystal used for analysis.

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINTE* (Bruker, 1998); data reduction: *SAINTE*; program(s) used to solve

structure: *SIR92* (Altomare *et al.*, 1994); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3* (Farrugia, 1997); software used to prepare material for publication: *SHELXL97*.

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