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Key indicators

Single-crystal X-ray study
T = 298 K
Mean $\sigma(\text{C}-\text{C}) = 0.009 \text{ \AA}$
R factor = 0.033
wR factor = 0.046
Data-to-parameter ratio = 15.5

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

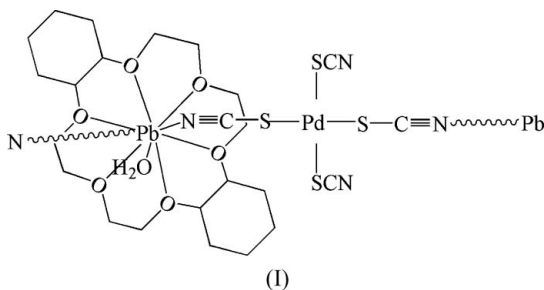
catena-Poly[[aqua(*cis-syn-cis*-dicyclohexyl-18-crown-6)lead(II)]- μ -thiocyanato- $\kappa^2\text{N}:\text{S}$ -[dithiocyanatopalladium(II)]- μ -thiocyanato- $\kappa^2\text{S}:\text{N}$]

The title polymeric complex, $[\text{PbPd}(\text{NCS})_4(\text{C}_{20}\text{H}_{36}\text{O}_6)(\text{H}_2\text{O})]_n$ or $\{[\text{Pb}(\text{DC18C6-A})(\text{H}_2\text{O})][\text{Pd}(\text{SCN})_4]\}_n$ (DC18C6-A = *cis-syn-cis*-dicyclohexyl-18-crown-6), has been synthesized by the reaction of dicyclohexyl-18-crown-6 with PdCl_2 and $\text{Pb}(\text{SCN})_2$. The SCN groups bridge $[\text{Pb}(\text{DC18C6-A})(\text{H}_2\text{O})]^{2+}$ complex cations and $[\text{Pd}(\text{SCN})_4]^{2-}$ complex anions to form infinite polymeric chains.

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Comment

In recent years, the synthesis and characterization of supramolecular architectures designed from metal-based coordination has expanded drastically (Reineke *et al.*, 1999; Ferey, 2001). This may be attributed largely to the potential applications of coordination polymers in many areas (Chen & Suslick, 1993; Lu, 2003). Our interest is focused on synthesizing some organic-inorganic hybrid crystalline solids containing crown ether cation building blocks (Dou *et al.*, 2004; Dong *et al.*, 2005). The title complex, (I), is reported here.



The molecular structure of (I) is shown in Fig. 1. The asymmetric unit consists of one $[\text{Pb}(\text{DC18C6-A})(\text{H}_2\text{O})]^{2+}$ complex cation (DC18C6-A = *cis-syn-cis*-dicyclohexyl-18-crown-6) and one $[\text{Pd}(\text{SCN})_4]^{2-}$ complex anion. In the

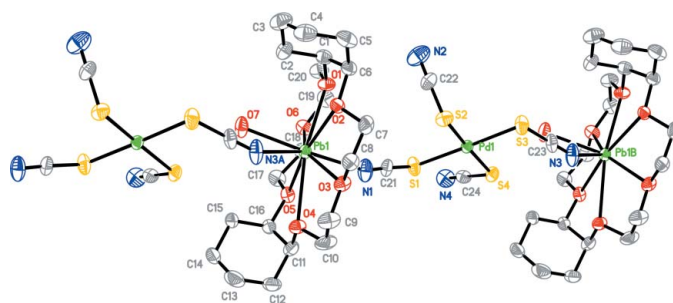


Figure 1
The molecular structure of (I), shown with 30% displacement ellipsoids, H atoms have been omitted for clarity [symmetry codes: (A) $x, y + 1, z$; (B) $x, y - 1, z$].

[Pd(SCN)₄]²⁻ complex anion, the Pd atom is coordinated by four S atoms from SCN groups and exhibits distorted square-planar coordination geometry. The bond lengths of Pd–S, S–C and C≡N are in the ranges 2.3082 (16)–2.3347 (17), 1.648 (6)–1.678 (7) and 1.128 (6)–1.160 (7) Å, respectively, which are consistent with the corresponding values found in the complex [Na(B18C6)]₂[Pd(SCN)₄](H₂O) (Li *et al.*, 2003). In the [Pb(DC18C6-A)(H₂O)]²⁺ complex cation, the Pb atom lies in the crown ether and is coordinated by nine atoms, six O atoms of a crown ether, one O atom from a water molecule and two N atoms from two [Pd(SCN)₄]²⁻ complex anions. The distances of Pb–O(crown ether) are in the range 2.666 (4)–2.839 (4) Å. The Pb1–N1 and Pb1–N3A bond lengths [symmetry code: (A: *x*, *y* + 1, *z*)] are 2.542 (5) and 2.927 (5) Å, respectively. Similar values can be found in the literature (Bailey *et al.*, 1983; Engelhardt *et al.*, 1989; Metz & Weiss, 1974). As shown in Fig. 1, the [Pb(DC18C6-A)(H₂O)]²⁺ complex cations and [Pd(SCN)₄]²⁻ complex anions are assembled into an infinite chain *via* SCN groups.

The coordinated water molecule forms two hydrogen bonds, *viz.* an intrachain O–H...S and an interchain O–H...N hydrogen bond (Table 1).

Experimental

All chemicals were commercially available and were used without further purification. The microanalytical data were obtained from a Perkin–Elmer 2400 II analyser. The title complex was prepared by adding an aqueous mixture (10 ml) of PdCl₂ (0.25 mmol) and Pb(SCN)₂ (5 mmol) to a solution of dicyclohexyl-18-crown-6 (1 mmol) in 1,2-dichloroethane (10 ml). The reaction mixture was stirred for 3 h at room temperature. The organic phase was separated from the reaction solution. Crystals of (I) were obtained from a 4:1 diethyl ether–1,2-dichloroethane solution over a period of 20 d.

Crystal data

[PbPd(NCS) ₄ (C ₂₀ H ₃₆ O ₆)(H ₂ O)]	<i>D_x</i> = 1.866 Mg m ⁻³
<i>M_r</i> = 936.41	Mo <i>K</i> α radiation
Monoclinic, <i>P</i> 2 ₁ / <i>n</i>	Cell parameters from 4334 reflections
<i>a</i> = 14.177 (5) Å	<i>θ</i> = 1.9–25.0°
<i>b</i> = 12.967 (4) Å	<i>μ</i> = 5.88 mm ⁻¹
<i>c</i> = 18.885 (6) Å	<i>T</i> = 298 (2) K
<i>β</i> = 106.237 (5)°	Block, red
<i>V</i> = 3333.4 (19) Å ³	0.25 × 0.23 × 0.18 mm
<i>Z</i> = 4	

Data collection

Bruker SMART CCD area-detector diffractometer	5836 independent reflections
<i>φ</i> and <i>ω</i> scans	3801 reflections with <i>I</i> > 2σ(<i>I</i>)
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)	<i>R</i> _{int} = 0.052
<i>T</i> _{min} = 0.241, <i>T</i> _{max} = 0.259	<i>θ</i> _{max} = 25.0°
17097 measured reflections	<i>h</i> = –15 → 16
	<i>k</i> = –15 → 15
	<i>l</i> = –20 → 22

Refinement

Refinement on <i>F</i> ²	H atoms treated by a mixture of independent and constrained refinement
<i>R</i> [<i>F</i> ² > 2σ(<i>F</i> ²)] = 0.033	<i>w</i> = 1/[σ ² (<i>F</i> _o ²) + (0.0003 <i>P</i>) ²]
<i>wR</i> (<i>F</i> ²) = 0.046	where <i>P</i> = (<i>F</i> _o ² + 2 <i>F</i> _c ²)/3
<i>S</i> = 0.95	(Δσ) _{max} = 0.001
5836 reflections	Δρ _{max} = 1.37 e Å ⁻³
376 parameters	Δρ _{min} = –0.95 e Å ⁻³

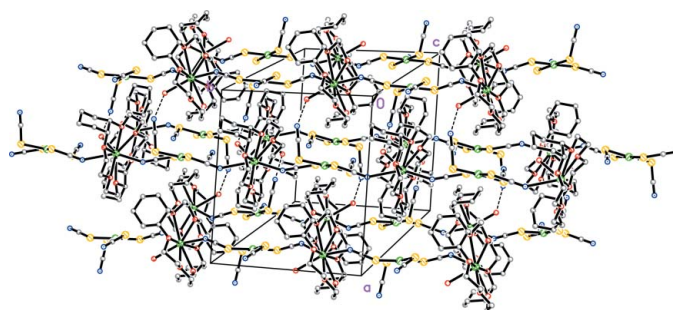


Figure 2
The unit-cell packing of (I), with dashed lines indicating the hydrogen bonds.

Table 1

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>
O7–H21...N2 ⁱ	0.90 (3)	1.93 (3)	2.805 (7)	165 (6)
O7–H22...S4 ⁱⁱ	0.87 (2)	2.54 (3)	3.349 (4)	154 (5)

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

Aqua H atoms were located in a difference map, the O–H distances were restrained to 0.88 (2) Å and the *U*_{iso}(H) values were fixed at 0.08 Å². Other H atoms were placed in idealized positions and constrained to ride on their parent atoms, with C–H = 0.97 Å and *U*_{iso}(H) = 1.2*U*_{eq}(C). The highest electron-density difference peak was located 1.03 Å from atom Pd1.

Data collection: SMART (Siemens, 1996); cell refinement: SAINT (Siemens, 1996); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997a); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997a); molecular graphics: SHELXTL (Sheldrick, 1997b); software used to prepare material for publication: SHELXTL.

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