

Tetrakis(thiourea-κS)palladium(II) dithiocyanate

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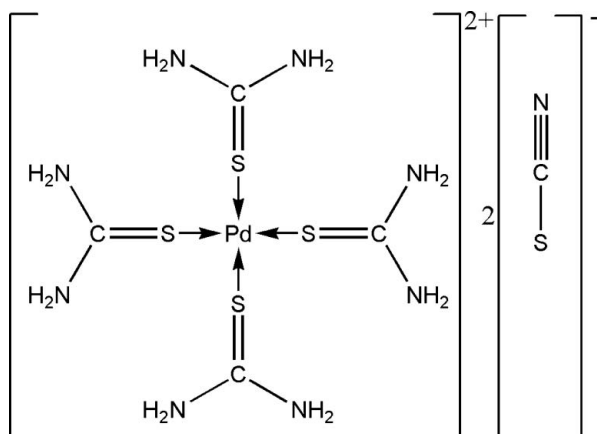
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Key indicators: single-crystal X-ray study; $T = 123$ K; mean $\sigma(\text{N}-\text{C}) = 0.003$ Å; R factor = 0.022; wR factor = 0.043; data-to-parameter ratio = 20.0.

The title compound, $[\text{Pd}(\text{CH}_4\text{N}_2\text{S})_4](\text{SCN})_2$, consists of complex $[\text{Pd}(\text{TU})_4]^{2+}$ [TU = thiourea, $\text{SC}(\text{NH}_2)_2$] cations and thiocyanate counter-anions. The Pd^{II} cation is situated on an inversion centre and exhibits an almost square-planar coordination by the S atoms of the TU ligands. The complex cations are connected through the thiocyanate ions *via* $\text{N}-\text{H}\cdots\text{N}$ [2.922 (3)–3.056 (3) Å] and $\text{N}-\text{H}\cdots\text{S}$ [3.369 (2)–3.645 (2) Å] hydrogen bonds.

Related literature

For the coordination chemistry of thiones and thionates, and for biomolecules possessing thioamido binding sites, see: Akrivos (2001); Raper (1996); Cusumano *et al.* (2005). For other structures listed in the Cambridge Structural Database (Allen, 2002) that contain transition metals and thiourea ligands, see: Bott *et al.* (1998); Dupa & Krebs (1973); Gale *et al.* (2006); Hunt *et al.* (1979); Taylor *et al.* (1974).



Experimental

Crystal data

$[\text{Pd}(\text{CH}_4\text{N}_2\text{S})_4](\text{SCN})_2$
 $M_r = 527.05$
 Monoclinic, $P2_1/c$
 $a = 8.136$ (3) Å
 $b = 12.966$ (5) Å
 $c = 8.810$ (3) Å
 $\beta = 91.12$ (5)°
 $V = 929.3$ (6) Å³
 $Z = 2$
 Mo $K\alpha$ radiation
 $\mu = 1.69$ mm⁻¹
 $T = 123$ (2) K
 $0.30 \times 0.25 \times 0.22$ mm

Data collection

Rigaku/MSC Mercury CCD diffractometer
 Absorption correction: integration (*NUMABS*; Higashi, 1999)
 $T_{\text{min}} = 0.632$, $T_{\text{max}} = 0.708$
 7301 measured reflections
 2117 independent reflections
 2040 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.025$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.022$
 $wR(F^2) = 0.043$
 $S = 1.35$
 2117 reflections
 106 parameters
 H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.66$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.48$ e Å⁻³

Table 1

Selected geometric parameters (Å, °).

Pd1–S2	2.3302 (11)	Pd1–S1	2.3448 (8)
S2–Pd1–S2 ⁱ	180	S2 ⁱ –Pd1–S1	92.14 (3)
S2–Pd1–S1	87.86 (3)		

Symmetry code: (i) $-x, -y + 1, -z + 1$.

Table 2

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
N1–H1B ⁱⁱ ⋯S3	0.88	2.58	3.369 (2)	150
N1–H1A ⁱⁱ ⋯S2 ⁱⁱⁱ	0.88	2.60	3.466 (2)	166
N2–H2A ⁱⁱ ⋯S3 ⁱⁱⁱ	0.88	2.78	3.615 (2)	158
N2–H2B ⁱⁱ ⋯N5 ^{iv}	0.88	2.04	2.922 (3)	178
N3–H3B ⁱⁱ ⋯S3	0.88	2.82	3.645 (2)	157
N3–H3A ⁱⁱ ⋯S1 ^v	0.88	2.73	3.531 (2)	153
N4–H4B ⁱⁱ ⋯S3 ^{vi}	0.88	2.61	3.482 (2)	173
N4–H4A ⁱⁱ ⋯N5 ^{vii}	0.88	2.50	3.056 (3)	121

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

Data collection: *CrystalClear* (Molecular Structure Corporation & Rigaku, 2001); cell refinement: *CrystalClear*; data reduction: *TEXSAN* (Molecular Structure Corporation & Rigaku, 2004); program(s) used to solve structure: *SIR97* (Altomare *et al.*, 1999); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEPII* (Johnson, 1976); software used to prepare material for publication: *SHELXL97* and *TEXSAN*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: WM2176).

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supporting information

Acta Cryst. (2008). E64, m698–m699 [doi:10.1107/S160053680801088X]

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S1. Comment

Thiourea (TU), $\text{SC}(\text{NH}_2)_2$, is a simple ambidentate ligand capable of binding to transition metals *via* the sulfur or the nitrogen atoms. Complex formation with such ligands provides model systems for the interaction of naturally occurring biomolecules possessing thioamido binding sites (Akrivos, 2001; Raper, 1996; Cusumano *et al.*, 2005). The ability of TU to form stable adducts with a variety of transition metals, *e.g.* Cu, Ag, Au and Pt, is well established. The crystal structures of several such complexes have been determined (Bott *et al.*, 1998; Gale *et al.*, 2006). These studies demonstrate that TU can act both as a terminal ligand in monomeric complexes (Hunt *et al.*, 1979), or as a bridging ligand in polymeric complexes (Taylor *et al.*, 1974). In order to investigate other transition metal complexes of thiourea, we report here the crystal structure of a monomeric complex, *viz.* $[\text{Pd}(\text{SC}(\text{NH}_2)_2)_4](\text{SCN})_2$, (I).

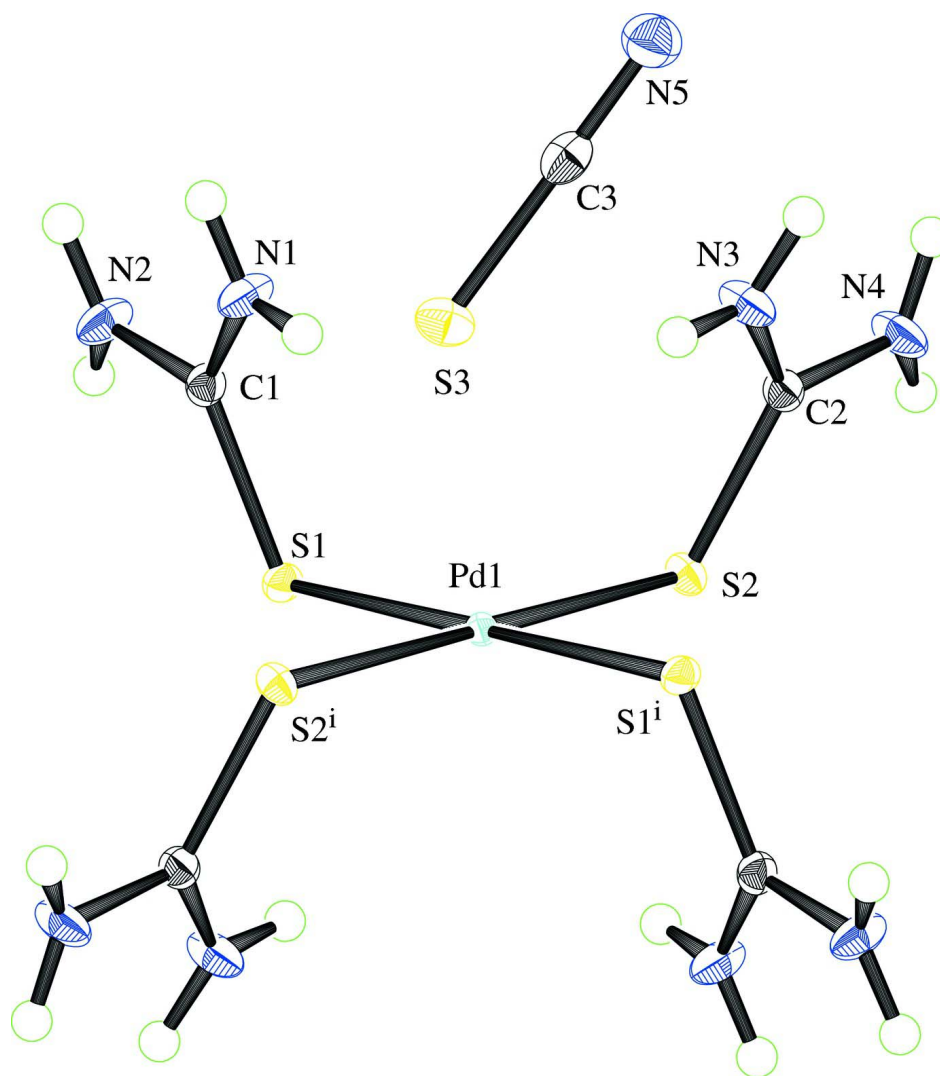
The crystal structure of (I) is composed of complex $[\text{Pd}(\text{TU})_4]^{+2}$ cations and thiocyanate counter anions. The Pd^{2+} ion is situated on an inversion centre and, as expected for a d^8 system, has an almost square planar environment with *cis* angles (S—Pd—S) ranging from 87.87 (2) to 92.13 (2)°, and *trans* angles (S—Pd—S) of 180.0°. The TU ligands are coordinated to Pd^{II} at almost equal distances. The Pd—S bond lengths of 2.3302 (8) and 2.3448 (7) Å (Table 1) are comparable to those of similar compounds reported in the literature (Gale *et al.*, 2006). In the cationic complex, TU ligands behave as S—donors and all four ligands are binding in a terminal mode. Therefore no bridging of metal centers are found as it is observed in some other metal-thiourea compounds, for example, $[\text{Cu}_4(\text{TU})_7(\text{SO}_4)_2]\text{NO}_3$ (Bott *et al.*, 1998) and $[\text{Ag}_2(\text{TU})_6](\text{ClO}_4)_2$ (Dupa & Krebs, 1973). The C—S and C—N bond lengths of 1.723 (2) Å and 1.326 (3) Å, respectively, agree with those of coordinated thiourea molecules reported in the Cambridge Crystallographic database (Allen, 2002). In the crystal structure, the building units are connected *via* hydrogen bonds of the type N—H \cdots N [2.922 (3)–3.058 (3) Å] and N—H \cdots S [3.370 (2)–3.646 (2) Å] (see Table 2).

S2. Experimental

Crystals of (I) were obtained by adding 4 equivalents of thiourea in 15 ml methanol to a solution of $\text{K}_2[\text{PdCl}_2]$ (0.326 g) in 15 ml of water and stirring for one h. The resulting orange solution was kept after filtration at room temperature for three d. Orange crystals of (I) were obtained on slow evaporation. The counter anion SCN^- has apparently been introduced due to impurities (presumably KSCN), that were present in thiourea.

S3. Refinement

The H atoms attached to the N atoms were placed in idealized positions and refined with a N—H distance of 0.88 Å and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{N})$.

**Figure 1**

Molecular structure of (I) showing the atom-labelling scheme and displacement ellipsoids at the 30% probability level. Unlabelled atoms and atoms labelled by superscript i) are related by the symmetry operator i) 1-x, y, z.

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Crystal data

$[\text{Pd}(\text{CH}_4\text{N}_2\text{S})_4](\text{SCN})_2$

$M_r = 527.05$

Monoclinic, $P2_1/c$

Hall symbol: -P 2ybc

$a = 8.136(3) \text{ \AA}$

$b = 12.966(5) \text{ \AA}$

$c = 8.810(3) \text{ \AA}$

$\beta = 91.12(5)^\circ$

$V = 929.3(6) \text{ \AA}^3$

$Z = 2$

$F(000) = 528$

$D_x = 1.884 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation, $\lambda = 0.71070 \text{ \AA}$

Cell parameters from 3103 reflections

$\theta = 3.4\text{--}27.5^\circ$

$\mu = 1.69 \text{ mm}^{-1}$

$T = 123 \text{ K}$

Prism, orange

$0.30 \times 0.25 \times 0.22 \text{ mm}$

Data collection

Rigaku/MS C Mercury CCD diffractometer	7301 measured reflections
Radiation source: fine-focus sealed tube	2117 independent reflections
Graphite monochromator	2040 reflections with $I > 2\sigma(I)$
ω scans	$R_{\text{int}} = 0.025$
Absorption correction: integration (NUMABS; Higashi, 1999)	$\theta_{\text{max}} = 27.5^\circ$, $\theta_{\text{min}} = 3.9^\circ$
$T_{\text{min}} = 0.632$, $T_{\text{max}} = 0.708$	$h = -8 \rightarrow 10$
	$k = -16 \rightarrow 13$
	$l = -11 \rightarrow 11$

Refinement

Refinement on F^2	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.022$	H-atom parameters constrained
$wR(F^2) = 0.043$	$w = 1/[\sigma^2(F_o^2) + 0.6382P]$
$S = 1.35$	where $P = (F_o^2 + 2F_c^2)/3$
2117 reflections	$(\Delta/\sigma)_{\text{max}} = 0.001$
106 parameters	$\Delta\rho_{\text{max}} = 0.66 \text{ e } \text{\AA}^{-3}$
0 restraints	$\Delta\rho_{\text{min}} = -0.48 \text{ e } \text{\AA}^{-3}$
Primary atom site location: structure-invariant direct methods	

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	$U_{\text{iso}}^*/U_{\text{eq}}$
Pd1	0.0000	0.5000	0.5000	0.00900 (6)
S1	-0.19061 (6)	0.37251 (3)	0.56476 (5)	0.01265 (10)
C1	-0.1354 (2)	0.26304 (14)	0.4664 (2)	0.0140 (4)
N1	-0.0189 (2)	0.26297 (13)	0.36425 (19)	0.0197 (4)
H1A	0.0068	0.2054	0.3175	0.024*
H1B	0.0332	0.3205	0.3429	0.024*
N2	-0.2134 (2)	0.17612 (13)	0.49809 (19)	0.0199 (4)
H2A	-0.1874	0.1187	0.4511	0.024*
H2B	-0.2911	0.1759	0.5661	0.024*
S2	0.13159 (6)	0.47003 (4)	0.73312 (5)	0.01300 (10)
C2	0.3204 (2)	0.41505 (14)	0.7012 (2)	0.0141 (4)
N3	0.3774 (2)	0.39827 (14)	0.56415 (18)	0.0201 (4)
H3A	0.4743	0.3693	0.5534	0.024*
H3B	0.3184	0.4160	0.4835	0.024*
N4	0.4105 (2)	0.38793 (14)	0.82155 (19)	0.0219 (4)
H4A	0.5073	0.3591	0.8096	0.026*

H4B	0.3736	0.3988	0.9134	0.026*
S3	0.22933 (7)	0.42269 (4)	0.17265 (6)	0.02044 (12)
C3	0.4076 (3)	0.36611 (15)	0.2055 (2)	0.0186 (4)
N5	0.5334 (2)	0.32525 (15)	0.2287 (2)	0.0260 (4)

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Pd1	0.00878 (10)	0.00771 (9)	0.01049 (9)	0.00000 (7)	-0.00061 (7)	0.00047 (7)
S1	0.0121 (2)	0.0103 (2)	0.0156 (2)	-0.00127 (17)	0.00147 (17)	-0.00054 (16)
C1	0.0152 (10)	0.0128 (9)	0.0137 (9)	-0.0004 (7)	-0.0027 (7)	0.0001 (7)
N1	0.0228 (10)	0.0129 (8)	0.0238 (9)	-0.0030 (7)	0.0077 (7)	-0.0057 (6)
N2	0.0248 (10)	0.0115 (8)	0.0236 (9)	-0.0039 (7)	0.0073 (7)	-0.0027 (7)
S2	0.0111 (2)	0.0163 (2)	0.0115 (2)	0.00134 (17)	-0.00052 (16)	0.00123 (16)
C2	0.0126 (9)	0.0124 (9)	0.0172 (9)	-0.0010 (7)	-0.0007 (7)	0.0011 (7)
N3	0.0166 (9)	0.0275 (9)	0.0162 (8)	0.0096 (7)	0.0005 (7)	0.0019 (7)
N4	0.0170 (9)	0.0309 (10)	0.0176 (8)	0.0108 (8)	-0.0040 (7)	0.0005 (7)
S3	0.0193 (3)	0.0202 (2)	0.0220 (2)	0.0026 (2)	0.0054 (2)	0.00404 (19)
C3	0.0235 (12)	0.0183 (10)	0.0144 (9)	-0.0049 (8)	0.0061 (8)	-0.0029 (7)
N5	0.0232 (11)	0.0256 (9)	0.0292 (10)	0.0018 (8)	0.0031 (8)	-0.0007 (8)

Geometric parameters (Å, °)

Pd1—S2	2.3302 (11)	N2—H2B	0.8800
Pd1—S2 ⁱ	2.3302 (11)	S2—C2	1.721 (2)
Pd1—S1	2.3448 (8)	C2—N3	1.320 (3)
Pd1—S1 ⁱ	2.3448 (8)	C2—N4	1.325 (3)
S1—C1	1.727 (2)	N3—H3A	0.8800
C1—N1	1.320 (3)	N3—H3B	0.8800
C1—N2	1.326 (3)	N4—H4A	0.8800
N1—H1A	0.8800	N4—H4B	0.8800
N1—H1B	0.8800	S3—C3	1.646 (2)
N2—H2A	0.8800	C3—N5	1.167 (3)
S2—Pd1—S2 ⁱ	180.0	C1—N2—H2B	120.0
S2—Pd1—S1	87.86 (3)	H2A—N2—H2B	120.0
S2 ⁱ —Pd1—S1	92.14 (3)	C2—S2—Pd1	108.72 (7)
S2—Pd1—S1 ⁱ	92.14 (3)	N3—C2—N4	119.29 (18)
S2 ⁱ —Pd1—S1 ⁱ	87.86 (3)	N3—C2—S2	123.27 (15)
S1—Pd1—S1 ⁱ	180.0	N4—C2—S2	117.44 (15)
C1—S1—Pd1	106.11 (7)	C2—N3—H3A	120.0
N1—C1—N2	119.74 (17)	C2—N3—H3B	120.0
N1—C1—S1	122.68 (15)	H3A—N3—H3B	120.0
N2—C1—S1	117.57 (15)	C2—N4—H4A	120.0
C1—N1—H1A	120.0	C2—N4—H4B	120.0
C1—N1—H1B	120.0	H4A—N4—H4B	120.0
H1A—N1—H1B	120.0	N5—C3—S3	179.5 (2)
C1—N2—H2A	120.0		

S2—Pd1—S1—C1	-101.78 (7)	S1—Pd1—S2—C2	112.12 (7)
S2 ⁱ —Pd1—S1—C1	78.22 (7)	S1 ⁱ —Pd1—S2—C2	-67.88 (7)
Pd1—S1—C1—N1	-6.95 (19)	Pd1—S2—C2—N3	2.44 (18)
Pd1—S1—C1—N2	172.15 (14)	Pd1—S2—C2—N4	-176.95 (14)

Symmetry code: (i) $-x, -y+1, -z+1$.

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>
N1—H1B...S3	0.88	2.58	3.369 (2)	150
N1—H1A...S2 ⁱⁱ	0.88	2.60	3.466 (2)	166
N2—H2A...S3 ⁱⁱⁱ	0.88	2.78	3.615 (2)	158
N2—H2B...N5 ^{iv}	0.88	2.04	2.922 (3)	178
N3—H3B...S3	0.88	2.82	3.645 (2)	157
N3—H3A...S1 ^v	0.88	2.73	3.531 (2)	153
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