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## Key indicators

Single-crystal X-ray study  
 $T = 298$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.004$  Å  
 $R$  factor = 0.023  
 $wR$  factor = 0.081  
Data-to-parameter ratio = 17.9For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.Dichlorobis[4-(methylsulfanyl)aniline- $\kappa$ N]-palladium(II)

In the title complex,  $[\text{PdCl}_2(\text{C}_7\text{H}_9\text{NS})_2]$ , the Pd atom lies on a center of inversion and is coordinated by two chloride anions and two N atoms from two benzenamine ligands to give rise to a square-planar coordination geometry for the Pd atom. A layer structure results from  $\text{N}-\text{H}\cdots\text{Cl}$  hydrogen bonds.

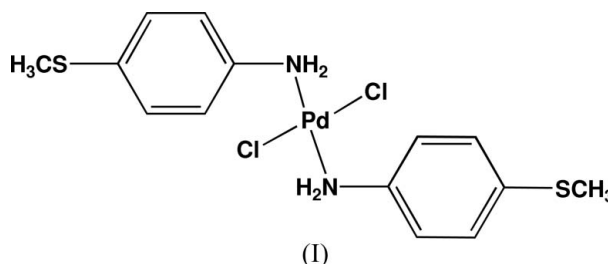
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## Comment

Palladium compounds have attracted much attention as a consequence of their applications in homogeneous and heterogeneous catalysis. For instance, these are active catalysts for the carbonylation of nitroarenes, aryl halides and alkynes (Bartolo *et al.*, 2001; Gaviño *et al.*, 2001; Gallo *et al.*, 1999). We report here the synthesis and the crystal structure of (I), a palladium dichloride adduct with 4-(methylthio)aniline, as a contribution to palladium chemistry.



The Pd atom lies on a center of inversion and is coordinated by two chloride anions and two N atoms from two substituted aniline ligands in a square-planar  $\text{PdCl}_2\text{N}_2$  geometry. In the crystal structure, molecules of (I) are linked *via*  $\text{N}-\text{H}\cdots\text{Cl}$  hydrogen bonds to form a layer structure.

## Experimental

The title compound was synthesized by the solvothermal reaction between 4-(methylthio)aniline and  $\text{PdCl}_2$  with a molar ratio of 2:1 in acetonitrile at 373 K for 10 h. After cooling to room temperature, orange crystals suitable for X-ray analysis were obtained.

## Crystal data

 $[\text{PdCl}_2(\text{C}_7\text{H}_9\text{NS})_2]$   
 $M_r = 455.72$   
Monoclinic,  $P2_1/c$   
 $a = 14.5222$  (19) Å  
 $b = 7.8998$  (10) Å  
 $c = 7.6175$  (10) Å  
 $\beta = 101.751$  (2)°  
 $V = 855.58$  (19) Å<sup>3</sup>  
 $Z = 2$  $D_x = 1.769$  Mg m<sup>-3</sup>  
Mo  $K\alpha$  radiation  
Cell parameters from 2975 reflections  
 $\theta = 2.4$ – $28.0^\circ$   
 $\mu = 1.63$  mm<sup>-1</sup>  
 $T = 298$  (2) K  
Prism, orange  
 $0.21 \times 0.17 \times 0.15$  mm

Data collection

Bruker SMART APEX area-detector diffractometer  
 $\varphi$  and  $\omega$  scans  
 Absorption correction: multi-scan (SADABS; Bruker, 2002)  
 $T_{\min} = 0.725$ ,  $T_{\max} = 0.792$   
 4730 measured reflections

1758 independent reflections  
 1661 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.018$   
 $\theta_{\text{max}} = 26.5^\circ$   
 $h = -18 \rightarrow 12$   
 $k = -9 \rightarrow 9$   
 $l = -9 \rightarrow 9$

Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.023$   
 $wR(F^2) = 0.081$   
 $S = 1.30$   
 1758 reflections  
 98 parameters  
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0466P)^2 + 0.0486P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\text{max}} = 0.001$   
 $\Delta\rho_{\text{max}} = 0.59 \text{ e \AA}^{-3}$   
 $\Delta\rho_{\text{min}} = -0.72 \text{ e \AA}^{-3}$

Table 1

Selected geometric parameters ( $\text{\AA}$ ,  $^\circ$ ).

Pd—N1	2.054 (2)	Pd—Cl	2.3060 (6)
N1—Pd—N1 <sup>i</sup>	180	N1—Pd—Cl <sup>i</sup>	90.91 (6)
N1—Pd—Cl	89.09 (6)	Cl—Pd—Cl <sup>i</sup>	180
Cl—Pd—N1—Cl	95.98 (16)	Cl <sup>i</sup> —Pd—N1—Cl	−84.02 (16)

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

Table 2

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N1—H1A $\cdots$ Cl <sup>ii</sup>	0.90	2.50	3.376 (2)	165
N1—H1B $\cdots$ Cl <sup>iii</sup>	0.90	2.50	3.321 (2)	152

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

All the H atoms were placed in geometrically idealized positions ( $C-H = 0.93 \text{ \AA}$  for the aromatic H atoms;  $C-H = 0.96 \text{ \AA}$  for the methyl H atoms;  $N-H = 0.90 \text{ \AA}$ ) and constrained to ride on their parent atoms, with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{aromatic C and N})$  or  $1.5U_{\text{eq}}(\text{methyl C})$ .

Data collection: SMART (Bruker, 2002); cell refinement: SAINT (Bruker, 2002); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics:

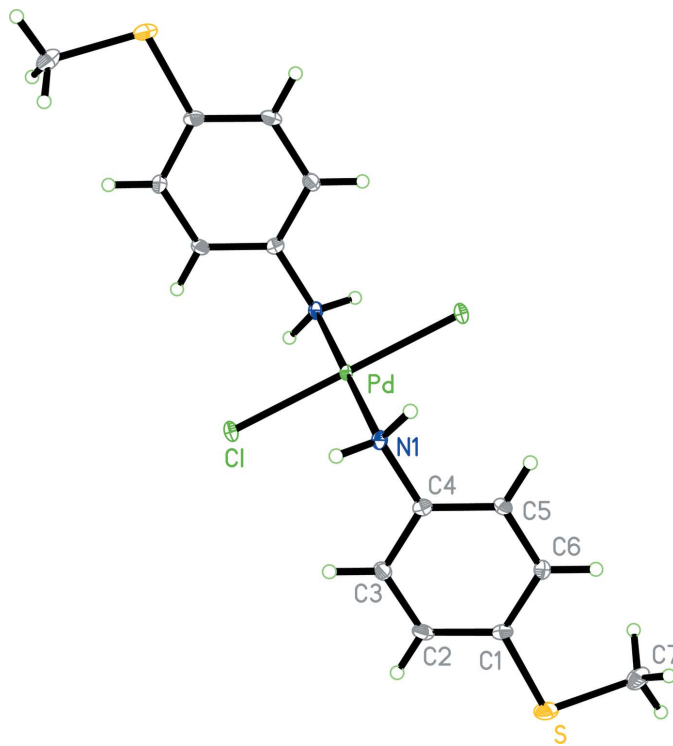


Figure 1

The structure of (I), showing 30% probability displacement ellipsoids and the atom-numbering scheme. Unlabeled atoms are related to labeled atoms by the symmetry operation  $(1 - x, 1 - y, 1 - z)$ .

SHELXTL (Bruker, 2002); software used to prepare material for publication: SHELXL97.

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