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## Key indicators

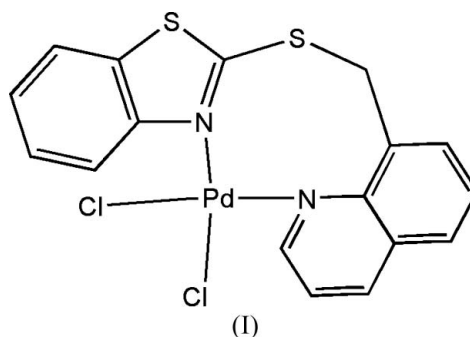
Single-crystal X-ray study  
 $T = 193$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.004$  Å  
 $R$  factor = 0.023  
 $wR$  factor = 0.059  
Data-to-parameter ratio = 14.5For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.Dichloro[8-(1,3-benzothiazol-2-ylsulfanylmethyl)-quinoline- $\kappa N, N'$ ]palladium(II)

In the mononuclear title complex,  $[\text{Pd}(\text{C}_{17}\text{H}_{12}\text{N}_2\text{S}_2)\text{Cl}_2]$ , the 8-(2-benzothiazolylsulfanylmethyl)quinoline ligand chelates to  $\text{PdCl}_2$  through its N atoms, conferring a square planar geometry on Pd. The angles around the Pd atom range from  $88.04(6)$  to  $92.76(3)^\circ$ .

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## Comment

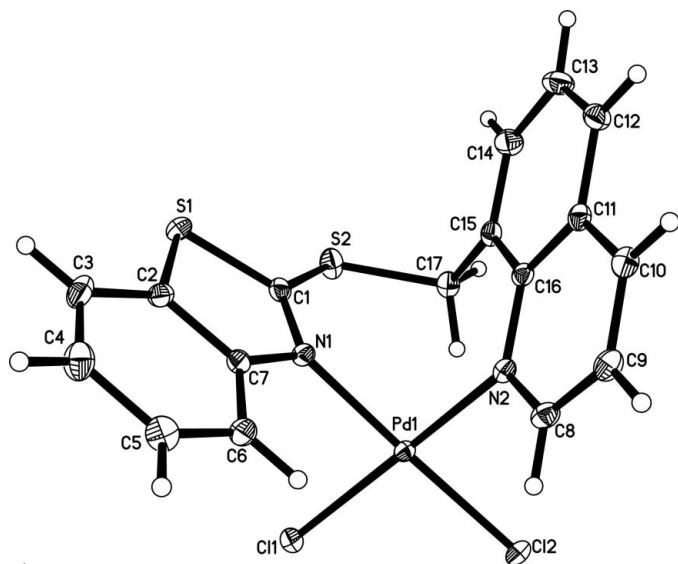
Square planar metal complexes have important applications in catalytic and bioinorganic systems (Fiallo & Garnier-Suillerot, 1986; Grundemann *et al.*, 2001; Trost *et al.*, 1995); for example, platinum(II) and palladium(II) complexes exhibit antitumour activity (Hay, 1987). The present study details the structure of a palladium dichloride adduct of a heterocyclic thioether ligand. A number of metal complexes of such ligands have been reported (Berry & Bebout, 2005; Song *et al.*, 2003; Zou *et al.*, 2004). The ligand used in this study is 8-(2-benzothiazolylsulfanylmethyl)quinoline.



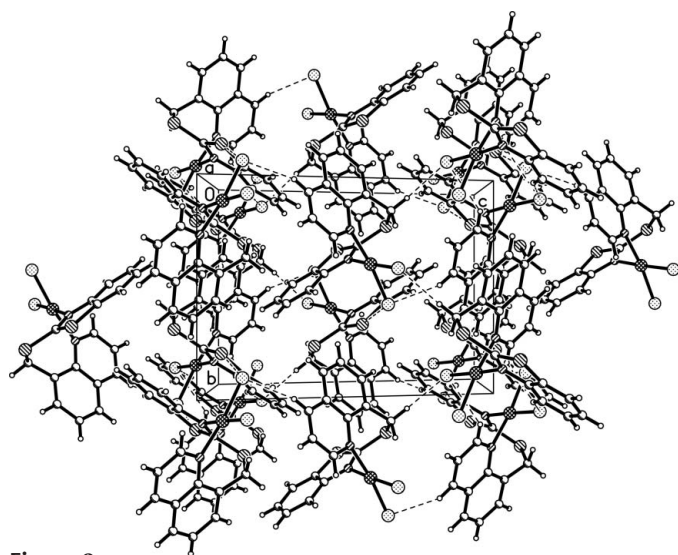
The title  $\text{PdCl}_2$  adduct, (I), is a mononuclear compound (Fig. 1 and Table 1), and the Pd exists in a square planar geometry formed by two Cl atoms and the two N atoms of the ligand. The quinoline and benzothiazole rings of the ligand are twisted by  $96.5(1)^\circ$ ; the Pd lies  $0.0387(1)$  Å from the square plane. The bond distances are within the ranges expected for square planar palladium (Al-Mandhary *et al.*, 2003; Buffin *et al.*, 2003; Kita *et al.*, 2002). The molecular packing in (I) is influenced by weak intermolecular C—H $\cdots$ Cl hydrogen bonds (Fig. 2 and Table 2).

## Experimental

The ligand (30.9 mg, 0.1 mmol) in  $\text{CHCl}_3$  (2 ml) was added to  $\text{PdCl}_2$  (17.8 mg, 0.1 mmol) dissolved in MeCN (20 ml). The mixture was stirred for 1 min, filtered and then set aside for the solvent to evaporate. Yield 34.5 mg (70%). Analysis found: C 42.10, H 2.46, N 5.76%; calculated for  $\text{C}_{17}\text{H}_{12}\text{Cl}_2\text{N}_2\text{PdS}_2$ : C 42.03, H 2.49, N 5.77%.



**Figure 1**  
View of (I), shown with 30% probability displacement ellipsoids and small spheres for the H atoms.



**Figure 2**  
The molecular packing of (I) viewed along the *a* axis. Dashed lines indicate hydrogen bonds.

#### Crystal data

[Pd(C<sub>17</sub>H<sub>12</sub>N<sub>2</sub>S<sub>2</sub>)Cl<sub>2</sub>]

*M<sub>r</sub>* = 485.71

Monoclinic, *P*2<sub>1</sub>/*c*

*a* = 12.417 (3) Å

*b* = 10.152 (3) Å

*c* = 14.541 (5) Å

β = 109.492 (5)°

*V* = 1728.0 (8) Å<sup>3</sup>

*Z* = 4

*D<sub>x</sub>* = 1.867 Mg m<sup>-3</sup>

Mo Kα radiation

Cell parameters from 7287

reflections

θ = 3.3–25.3°

μ = 1.63 mm<sup>-1</sup>

*T* = 193 (2) K

Block, orange

0.46 × 0.35 × 0.23 mm

#### Data collection

Rigaku Mercury diffractometer

ω scans

Absorption correction: multi-scan  
(Jacobson, 1998)

*T<sub>min</sub>* = 0.522, *T<sub>max</sub>* = 0.706

16474 measured reflections

3157 independent reflections

3034 reflections with *I* > 2σ(*I*)

*R<sub>int</sub>* = 0.023

θ<sub>max</sub> = 25.3°

*h* = -14 → 13

*k* = -12 → 12

*l* = -16 → 17

#### Refinement

Refinement on *F*<sup>2</sup>

*R* [*F*<sup>2</sup> > 2σ(*F*<sup>2</sup>)] = 0.023

*wR* (*F*<sup>2</sup>) = 0.059

*S* = 1.11

3157 reflections

218 parameters

H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.0245P)^2 + 2.5404P]$$

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

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

$$\Delta\rho_{\max} = 0.55 \text{ e } \text{Å}^{-3}$$

$$\Delta\rho_{\min} = -0.56 \text{ e } \text{Å}^{-3}$$

**Table 1**

Selected geometric parameters (Å, °).

Pd1–N1	2.035 (2)	Pd1–Cl2	2.2971 (8)
Pd1–N2	2.045 (2)	Pd1–Cl1	2.3007 (8)
N1–Pd1–N2	89.52 (8)	N1–Pd1–Cl1	88.04 (6)
N1–Pd1–Cl2	177.58 (6)	N2–Pd1–Cl1	173.08 (6)
N2–Pd1–Cl2	89.94 (6)	Cl2–Pd1–Cl1	92.76 (3)

**Table 2**

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>
C17–H17B···Cl2 <sup>i</sup>	0.99	2.75	3.667 (3)	154
C10–H10···Cl1 <sup>ii</sup>	0.95	2.71	3.466 (3)	137
C9–H9···Cl1 <sup>iii</sup>	0.95	2.94	3.658 (3)	133
C8–H8···Cl2 <sup>iii</sup>	0.95	2.84	3.575 (3)	135
C6–H6···Cl2 <sup>iii</sup>	0.95	2.73	3.571 (3)	148
C3–H3···Cl1 <sup>iv</sup>	0.95	2.92	3.644 (3)	134

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

H atoms were included in calculated positions and refined as riding on C, with C–H distances of 0.95 Å (aromatic H) and 0.99 Å (methylene H); *U*<sub>iso</sub>(H) = 1.2*U*<sub>eq</sub>(C).

Data collection: *CrystalClear* (Molecular Structure Corporation, 2000; Rigaku Corporation, 1999); cell refinement: *CrystalClear*; data reduction: *CrystalStructure* (Rigaku & Rigaku/MS, 2003); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 1998); software used to prepare material for publication: *SHELXTL*.

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