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Key indicators

Single-crystal X-ray study

 $T = 233$ KMean $\sigma(\text{C}-\text{C}) = 0.004$ Å R factor = 0.026 wR factor = 0.061

Data-to-parameter ratio = 15.7

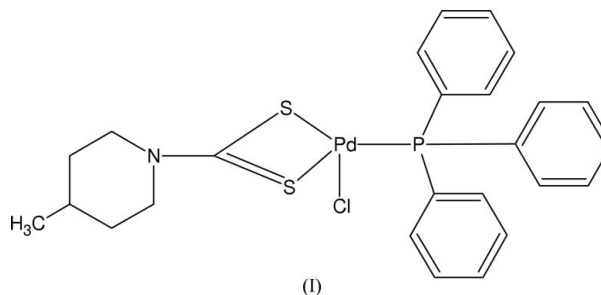
For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.Chloro(4-methylpiperidine-1-dithiocarbamato- $\kappa^2\text{S:S}'$)-
(triphenylphosphine- κP)palladium(II)In the title compound, $[\text{Pd}(\text{C}_7\text{H}_{12}\text{NS}_2)\text{Cl}(\text{C}_{18}\text{H}_{15}\text{P})]$, the Pd
atom is four-coordinate and exhibits a slightly distorted
square-planar geometry.

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Comment

Palladium(II) complexes with sulfur and phosphorus donor
ligands are of current interest due to their ability to sequester
the metal ion (Faraglia *et al.*, 2005), their antitumor activity
against leukemic cells (Mital *et al.*, 1989; Tiekink, 2002), and
their use as pesticides (Fackler, 2002) and antimicrobial agents
(Ronconi *et al.*, 2005), while palladium(II)-phosphine
complexes are also important from a catalytic point of view
(Crawforth *et al.*, 2005; Tsuji, 1995), *e.g.* $[\text{Pd}(\text{PPh}_3)_2(\text{CN})_2]$
(Hua *et al.*, 2001), $[\text{Pd}(\text{PPh}_3)_2\text{Cl}_2]$ (Nicholas, 1987).In the title compound, (I), the dithiocarbamate ligand acts
as a bidentate chelate, coordinating to Pd *via* both S atoms.
Atom S2 is *trans* to the chloro ligand and atom S1 *trans* to the
triphenylphosphine ligand (Fig. 1 and Table 1). The coordi-
nation geometry about the Pd atom is distorted square planar
and the deviation of the Pd1 atom from the mean plane
through the ligand donor atoms is only 0.0103 (4) Å.

Experimental

4-Methylpiperidine-1-dithiocarbamic acid (Vogel, 1968) dissolved
(0.2 g, 1.14 mmol) in CH_2Cl_2 (10 ml) was added to a suspension of
 $[\text{PdCl}_2(\text{PPh}_3)]$ (0.5 g, 1.14 mmol) (Kitano *et al.*, 1983) in CH_2Cl_2
(20 ml). The resulting solution was refluxed for 1 h. Yellow crystals
were obtained on slow evaporation of the solvent at room
temperature.

Crystal data

 $[\text{Pd}(\text{C}_7\text{H}_{12}\text{NS}_2)\text{Cl}(\text{C}_{18}\text{H}_{15}\text{P})]$ $M_r = 578.42$ Monoclinic, $P2_1/n$ $a = 10.1668$ (3) Å $b = 13.8325$ (4) Å $c = 17.8608$ (4) Å $\beta = 90.162$ (2)° $V = 2511.79$ (12) Å³ $Z = 4$ $D_x = 1.530$ Mg m⁻³Mo $K\alpha$ radiation

Cell parameters from 13105

reflections

 $\theta = 1.0$ – 26.0 ° $\mu = 1.09$ mm⁻¹ $T = 233$ (2) K

Prism, yellow

 $0.2 \times 0.12 \times 0.1$ mm

Data collection

Nonius KappaCCD diffractometer
 φ and ω scans
 Absorption correction: none
 13105 measured reflections
 4405 independent reflections
 3853 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.025$
 $\theta_{\text{max}} = 25.0^\circ$
 $h = -12 \rightarrow 12$
 $k = -16 \rightarrow 16$
 $l = -21 \rightarrow 21$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.026$
 $wR(F^2) = 0.061$
 $S = 1.05$
 4405 reflections
 280 parameters
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0201P)^2 + 2.004P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.002$
 $\Delta\rho_{\text{max}} = 0.39 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.55 \text{ e } \text{\AA}^{-3}$

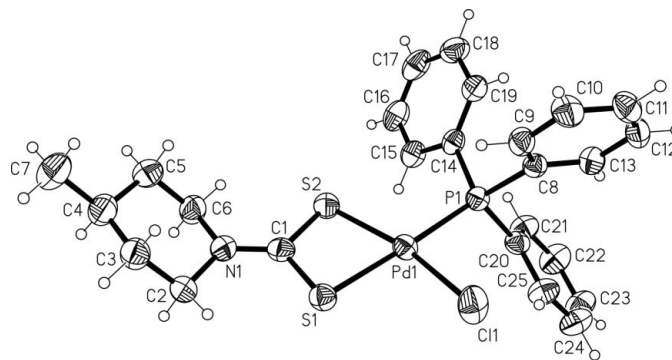


Figure 1

The structure of (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level.

Table 1

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

Pd1—P1	2.3009 (7)	S1—C1	1.718 (3)
Pd1—Cl1	2.3276 (7)	S2—C1	1.735 (3)
Pd1—S1	2.3274 (7)	N1—C1	1.313 (3)
Pd1—S2	2.2977 (7)		
S2—Pd1—P1	100.32 (2)	C1—S1—Pd1	87.06 (9)
S2—Pd1—S1	75.33 (2)	C1—S2—Pd1	87.61 (9)
P1—Pd1—S1	175.24 (2)	N1—C1—S1	125.1 (2)
S2—Pd1—Cl1	166.37 (3)	N1—C1—S2	124.99 (19)
P1—Pd1—Cl1	93.29 (2)	S1—C1—S2	109.86 (14)
S1—Pd1—Cl1	91.08 (3)		

H atoms were positioned geometrically ($\text{C—H} = 0.94\text{--}0.98 \text{ \AA}$) and refined as riding, with $U_{\text{iso}}(\text{H}) = 1.2$ or 1.5 times $U_{\text{eq}}(\text{C})$.

Data collection: *COLLECT* (Nonius, 1998); cell refinement: *SCALEPACK* (Otwinowski & Minor, 1997); data reduction: *DENZO* (Otwinowski & Minor, 1997) and *SCALEPACK*; 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: *SHELXL97*.

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