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Key indicators

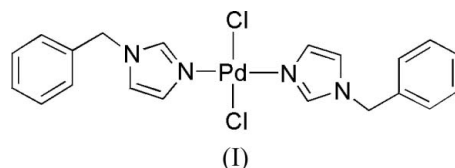
Single-crystal X-ray study
 $T = 150$ K
Mean $\sigma(\text{C}-\text{C}) = 0.002$ Å
 R factor = 0.018
 wR factor = 0.048
Data-to-parameter ratio = 19.3For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.*trans*-Bis(1-benzylimidazole)dichloropalladium(II)

The structure of the title compound, $[\text{PdCl}_2(\text{C}_{10}\text{H}_{10}\text{N}_2)_2]$, has the palladium center situated on a center of inversion. The imidazole group is twisted from the square coordination plane with a $\text{C}-\text{N}-\text{Pd}-\text{Cl}$ torsion angle of $31.93(12)^\circ$; this is different from the angle reported in closely related structures.

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Comment

Our group is interested in the preparation of palladium complexes of *N*-heterocyclic carbene (NHC) ligands (Lee *et al.*, 2004). A general synthetic method is *via* the *in situ* deprotonation of the corresponding imidazolium salt to generate the NHC ligand, which is then trapped by a suitable palladium precursor. In the course of preparing a palladium dichloride complex supported by a multidentate NHC ligand, we reacted the corresponding imidazolium salt and palladium dichloride. An unsuccessful attempt resulted in the decomposition product *trans*-bis(1-benzylimidazole)dichloropalladium(II), (I), which can be independently prepared by the reaction between 1-benzylimidazole and palladium(II) dichloride in dimethyl sulfoxide (DMSO).



Here, we present the structure of (I) (Fig. 1). It crystallizes in the monoclinic space group $P2_1/n$ with the palladium center situated at a center of inversion. The imidazole unit is twisted away from the square coordination plane, with a $\text{C}2-\text{N}2-$

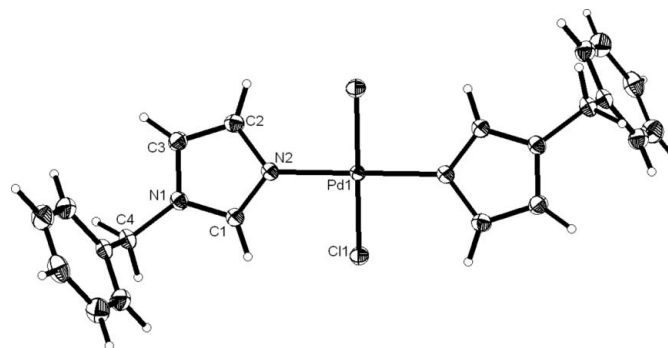


Figure 1

The structure of (I), showing 50% probability displacement ellipsoids for non-H atoms. H atoms are of arbitrary size. Unlabeled atoms are related to labeled atoms by $(1-x, -y, -z)$.

Pd1—C11 torsion angle of $31.93(12)^\circ$. The corresponding angles in a closely related structure with an *N-n*-propyl substituent on the imidazole ligand are, however, $19.0(3)$ and $23.0(3)^\circ$ (Ianelli *et al.*, 1985), whereas that in another complex with an *N*-vinyl substituent is $47.6(1)^\circ$ (Kurdziel & Glowiak, 2002).

Experimental

A mixture of 1-benzylimidazole (0.221 g, 1.40 mmol) and palladium(II) dichloride (0.124 g, 0.700 mmol) was heated at 363 K for 2 d in DMSO (10 ml). After cooling, the solvent was removed completely under vacuum to afford a yellow solid (yield 0.283 g, 82%). Crystals suitable for structural analysis were obtained by vapor diffusion of diethyl ether into a dimethylformamide solution containing the solid.

Crystal data

[PdCl ₂ (C ₁₀ H ₁₀ CIN ₂) ₂]	<i>Z</i> = 2
<i>M_r</i> = 493.70	<i>D_x</i> = 1.686 Mg m ⁻³
Monoclinic, <i>P</i> 2 ₁ / <i>n</i>	Mo <i>K</i> α radiation
<i>a</i> = 6.7632 (1) Å	<i>μ</i> = 1.24 mm ⁻¹
<i>b</i> = 20.6542 (4) Å	<i>T</i> = 150 (2) K
<i>c</i> = 7.5512 (1) Å	Prism, yellow
<i>β</i> = 112.749 (2)°	0.25 × 0.20 × 0.17 mm
<i>V</i> = 972.76 (3) Å ³	

Data collection

Bruker SMART APEXII diffractometer	9632 measured reflections
<i>ω</i> scans	2393 independent reflections
Absorption correction: multi-scan (SADABS; Sheldrick, 2003)	2170 reflections with <i>I</i> > 2σ
<i>T_{min}</i> = 0.747, <i>T_{max}</i> = 0.817	<i>R_{int}</i> = 0.018
	<i>θ_{max}</i> = 28.2°

Refinement

Refinement on <i>F</i> ²	$w = 1/[\sigma^2(F_o^2) + (0.0249P)^2 + 0.4092P]$
$R[F^2 > 2\sigma(F^2)] = 0.018$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.048$	$(\Delta/\sigma)_{\max} = 0.001$
<i>S</i> = 1.06	$\Delta\rho_{\max} = 0.47 \text{ e \AA}^{-3}$
2393 reflections	$\Delta\rho_{\min} = -0.47 \text{ e \AA}^{-3}$
124 parameters	
H-atom parameters constrained	

All H atoms were positioned geometrically and refined with a riding model, with *U_{iso}*(H) = 1.2*U_{eq}*(C) and C—H = 0.95–0.99 Å.

Data collection: APEX2 (Bruker, 2004); cell refinement: SAINT (Bruker, 2004); data reduction: SAINT; program(s) used to solve structure: SHELXTL (Sheldrick, 1998); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL.

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