metal-organic papers

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Key indicators

Single-crystal X-ray study T = 100 KMean σ (C–C) = 0.007 Å R factor = 0.048 wR factor = 0.118 Data-to-parameter ratio = 10.6

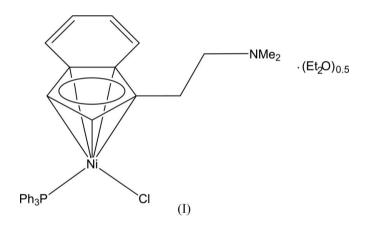
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

Chloro[η^5 -1-(dimethylaminoethyl)indenyl]-(triphenylphosphine)nickel(II) diethyl ether hemisolvate

The title complex, $[Ni(C_{13}H_{16}N)Cl(C_{18}H_{15}P)]\cdot 0.5C_4H_{10}O$, consists of an Ni^{II} center coordinated by a dimethylaminoethyl-substituted indenyl ligand, a triphenylphosphine ligand and a chloride ion. The asymmetric unit includes two very similar molecules that show only small conformational differences in the indenyl side chain. Received 30 June 2005 Accepted 4 July 2005 Online 9 July 2005

Comment

The crystallographic study of the title compound, (I), was undertaken during the course of our studies on substituted indenyl complexes. The asymmetric unit contains two independent complex molecules and a diethyl ether molecule. One of the two complex molecules is represented in Fig. 1. The other has the same structure with minor differences in the conformation of the side chain.



Crystal structures of two solvent-free polymorphs of the same complex have already been published by our group (Groux *et al.*, 2000; Groux & Zargarian, 2001). All three crystal structures display a similar coordination environment around the Ni atom and spacial orientations for the indenyl group, the amine side chain, and the phenyl rings of the triphenylphosphine ligand. The solvent-free materials show disorder in the side chain, but there is no evidence of such disorder in the present structure.

Experimental

The preparation of the main title complex has already been published (Groux *et al.*, 2000). The single crystals of (I) obtained in the present study were grown from a solution of the complex in diethyl ether/hexanes.

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Crystal data

$$\begin{split} & [\mathrm{Ni}(\mathrm{C}_{13}\mathrm{H}_{16}\mathrm{N})\mathrm{Cl}(\mathrm{C}_{18}\mathrm{H}_{15}\mathrm{P})] & \cdot \\ & 0.5\mathrm{C}_4\mathrm{H}_{10}\mathrm{O} \\ & M_r = 579.77 \\ & \mathrm{Triclinic}, \ P\overline{1} \\ & a = 12.7839 \ (5) \ \mathring{\mathrm{A}} \\ & b = 12.9973 \ (6) \ \mathring{\mathrm{A}} \\ & c = 18.0683 \ (8) \ \mathring{\mathrm{A}} \\ & \alpha = 100.725 \ (2)^{\circ} \\ & \beta = 98.363 \ (2)^{\circ} \\ & \gamma = 90.364 \ (2)^{\circ} \\ & V = 2916.6 \ (2) \ \mathring{\mathrm{A}}^3 \end{split}$$

Data collection

Bruker SMART2000 diffractometer ω scans Absorption correction: multi-scan (*SADABS*; Bruker, 1999) $T_{min} = 0.600, T_{max} = 0.770$ 36298 measured reflections 7196 independent reflections

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.048$ $wR(F^2) = 0.118$ S = 0.957196 reflections 682 parameters

Table 1

Selected bond lengths (Å).

Ni-C3	2.045 (4)	Ni1-C13	2.014 (4)
Ni-C2	2.068 (4)	Ni1-C12	2.059 (5)
Ni-C1	2.142 (4)	Ni1-C11	2.168 (4)
Ni-C3a	2.332 (5)	Ni1-C13a	2.306 (4)
Ni-C7a	2.366 (5)	Ni1-C17a	2.361 (4)
Ni-P	2.1758 (14)	Ni1-P1	2.1846 (14)
Ni-Cl	2.1976 (13)	Ni1-Cl1	2.1925 (13)

Z = 4

 $D_x = 1.320 \text{ Mg m}^{-3}$

Cell parameters from 5841

 $0.20 \times 0.15 \times 0.12 \text{ mm}$

4733 reflections with $I > 2\sigma(I)$

H-atom parameters constrained

 $w = 1/[\sigma^{2}(F_{o}^{2}) + (0.0598P)^{2}]$ where $P = (F_{o}^{2} + 2F_{c}^{2})/3$

Cu Ka radiation

reflections

 $\mu=2.50~\mathrm{mm}^{-1}$

T = 100 (2) K

Block, red

 $R_{\rm int} = 0.046$

 $\theta_{\rm max} = 55.2^{\circ}$

 $h = -13 \rightarrow 13$

 $k = -13 \rightarrow 13$

 $l = -19 \rightarrow 19$

 $(\Delta/\sigma)_{\rm max} = 0.009$

 $\Delta \rho_{\rm max} = 0.77 \ {\rm e} \ {\rm \AA}^{-3}$

 $\Delta \rho_{\rm min} = -0.46 \text{ e } \text{\AA}^{-3}$

 $\theta = 3.5 - 55.1^{\circ}$

The H atoms were placed in calculated positions (C-H = 0.93–0.98 Å) and refined as riding with the constraint $U_{\rm iso}({\rm H}) = 1.2U_{\rm eq}({\rm carrier})$ or $1.5U_{\rm eq}({\rm methyl \ carrier})$ applied.

Data collection: *SMART* (Bruker, 1999); cell refinement: *SAINT* (Bruker, 1999); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 1997); software used to prepare material for publication: *UdMX* (Maris, 2004).

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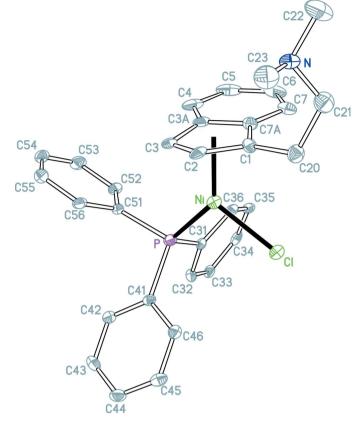


Figure 1

View of one of the Ni-containing molecules in (I), showing 30% displacement ellipsoids. H atoms have been omitted for clarity and the interaction between Ni and the centroid of the five-membered ring is shown. The second molecule has a very similar structure.

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