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# **Structure Reports**

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#### **Key indicators**

Single-crystal X-ray study T = 297 KMean  $\sigma(\text{C-C}) = 0.009 \text{ Å}$ Disorder in main residue R factor = 0.059 wR factor = 0.169 Data-to-parameter ratio = 13.9

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

# rac-[2,2'-Bis(1-ethylbenzimidazol-2-yl- $\kappa N^3$ )-biphenyl]dichloronickel(II)

The title compound,  $[NiCl_2(C_{30}H_{26}N_4)]$ , is a monoclinic (purple) racemic polymorph of a previously reported orthorhombic (blue) form of the same compound. The complex exhibits pseudo-tetrahedral coordination geometry around the nickel(II) metal center. This geometry is enforced by the geometrically constraining ligand 2,2'-bis(1-ethylbenzimidazol-2-yl)biphenyl.

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#### Comment

Bis(2-benzimidazol-2-yl) species bridged connecting moieties have attracted considerable interest (Stibrany et al., 2003b). In the present study, we have examined a racemic polymorphic form of a 2,2'-biphenyl-bridged bis(2benzimidazol-2-yl) species complexed to nickel(II). When the geometrically constraining ligand 2,2'-bis(1-ethylbenzimidazol-2-yl)biphenyl, (I) (Stibrany et al., 2003a) is complexed to nickel(II) dichloride, a pseudo-tetrahedral coordination environment is adopted by the nickel metal center, (II). These nickel(II) pre-catalysts can be activated with methylaluminoxane to oligomerize ethylene (Stibrany et al., 2002) and to polymerize norbornene (Patil et al., 2003). The analogous copper(II) complexes can be activated with methylaluminoxane to polymerize ethylene and acrylates (Stibrany, Schulz et al., 2003).

The monoclinic title compound, (II)<sub>m</sub>, crystallizes as a racemate. Chirality is necessarily introduced by the 2,2'-bi-phenyl bridge. Facile interconversion of the biphenyl bridge in the free ligand is expected due to the low rotational barrier offered by the 6,6'-proton substitution. The crystal chosen of the polymorphic orthorhombic form (II)<sub>o</sub> was crystallographically determined to be the R enantiomer (Stibrany, Matturro et al., 2003). It is not yet known if (II)<sub>m</sub> is the result of a solvent-induced racemization of R-(II)<sub>o</sub> or if (II)<sub>o</sub> exists as a mixture of R and S crystals. The nickel coordination varies only slightly in the two polymorphic forms. The nickel coordination parameters for (II)<sub>m</sub> are Ni-N = 2.000 (3)-2.005 (3) Å and Ni-Cl = 2.2310 (15)-2.2405 Å, and the angles are 98.55 (11)-125.90 (6)°. In (II)<sub>o</sub>, the corresponding ranges are 1.999 (8)-2.008 (7) Å, 2.226 (2)-2.233 (3) Å and 101.0 (2)-

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© 2004 International Union of Crystallography Printed in Great Britain – all rights reserved  $123.43 (11)^{\circ}$ . In (II)<sub>m</sub>, the phenyl-phenyl dihedral angle is  $81.6 (2)^{\circ}$ , compared with  $83.9 (3)^{\circ}$  in (II)<sub>o</sub>.

Occupation of the calculated solvent-accessible void of 37  $\text{Å}^3$  for (II)<sub>m</sub> is unlikely. The complex was prepared in the presence of triethylorthoformate to ensure the removal of any adventitious water. The measured density of 1.34 (1) Mg m<sup>-3</sup> is within experimental error of the value (1.359 Mg m<sup>-3</sup>) calculated from the structure. The calculated density is 1.402 Mg m<sup>-3</sup> if a water molecule occupies the void. The measured density of the blue form was found to be 1.37 (1) Mg m<sup>-3</sup> which compares well with the calculated density of 1.372 Mg m<sup>-3</sup>. The blue form,  $(II)_o$ , does not have a void large enough to accommodate a solvent molecule. This accounts for the slightly higher density. The measured density of the bulk blue form seems to indicate a uniform density. This indicates that either the S enantiomer has nearly the same density as the R enantiomer or the blue form exists entirely as the R enantiomer. In either case, a second method of separation of blue and purple forms is provided by flotation.

## **Experimental**

The blue orthorhombic crystalline form of the title compound, (II), was obtained as previously described in Stibrany et al. (2002). This solid was dissolved in a mixture of acetonitrile and triethylorthoformate with gentle reflux. The blue-purple solution was then sealed in a jar to allow diethyl ether diffusion. Overnight, a mixture of blue and purple crystals was obtained which could be physically separated.

### Crystal data

$[NiCl_2(C_{30}H_{26}N_4)]$	$D_m$ measured by flotation in
$M_r = 572.15$	CCl <sub>4</sub> /cyclohexane
Monoclinic, $P2_{\downarrow}/n$	Mo $K\alpha$ radiation
a = 9.1464 (5)  Å	Cell parameters from 765
b = 18.6955 (10) Å	reflections
c = 16.4401 (10)  Å	$\theta = 2.5 - 21.3^{\circ}$
$\beta = 95.806 (4)^{\circ}$	$\mu = 0.91 \text{ mm}^{-1}$
$V = 2796.8 (3) \text{ Å}^3$	T = 297 (1)  K
Z = 4	Rod, purple
$D_x = 1.359 \text{ Mg m}^{-3}$	$0.46 \times 0.12 \times 0.04 \text{ mm}$
$D_m = 1.34 (1) \text{ Mg m}^{-3}$	

### Data collection

Siemens P3 diffractometer with a	4918 independent reflections
Bruker SMART APEX CCD	3033 reflections with $I > 2\sigma(I)$
area-detector	$R_{\rm int} = 0.071$
$\varphi$ and $\omega$ scans	$\theta_{\rm max} = 25.0^{\circ}$
Absorption correction: multi-scan	$h = -10 \rightarrow 10$
(SADABS; Blessing, 1995)	$k = -22 \rightarrow 22$
$T_{\min} = 0.787, T_{\max} = 0.960$	$l = -15 \rightarrow 19$
20111 measured reflections	

### Refinement

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Refinement on $F^2$	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.059$	$w = 1/[\sigma^{2}(F_{o}^{2}) + (0.0957P)^{2}]$
$wR(F^2) = 0.169$	where $P = (F_o^2 + 2F_c^2)/3$
S = 1.00	$(\Delta/\sigma)_{\rm max} < 0.001$
4918 reflections	$\Delta \rho_{\text{max}} = 0.63 \text{ e Å}^{-3}$
353 parameters	$\Delta \rho_{\text{min}} = -0.43 \text{ e Å}^{-3}$

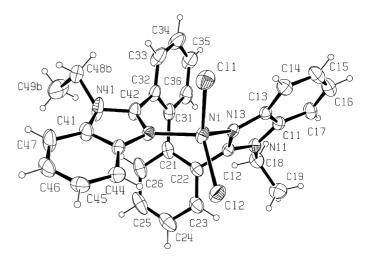


Figure 1 The molecular structure of  $(II)_m$ , showing 25% probability displacement ellipsoids. Only the minor component of the disordered ethyl group is shown.

One of the ethyl groups C48-C49 was found to be disordered over two sites. The disorder also resulted in some unusually large deviations in the near neighbor displacement parameters.

Data collection: SMART (Bruker, 2000); cell refinement: SMART; data reduction: SAINT (Bruker, 2000); program(s) used to solve structure: SHELXS97 (Sheldrick, 1990); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEPIII (Burnett & Johnson, 1996) and ORTEP32 (Farrugia, 1997); software used to prepare material for publication: SHELXTL (Bruker, 2000).

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