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Key indicators

Single-crystal X-ray study T = 150 K Mean σ (C–C) = 0.003 Å R factor = 0.039 wR factor = 0.096 Data-to-parameter ratio = 9.9

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

(*S*,*S*)-*N*,*N*'-Dicyanoethyl-1,2-(2-hydroxyphenyl)ethylenediamine

In the title molecule {systematic name: 3,3'-[1,2-bis(2-hydroxyphenyl)ethane-1,2-diyldiimino]propanenitrile}, $C_{20}H_{22}N_4O_2$, there are three different types of intramolecular hydrogen bonds, one of each of the types $N-H\cdots O$ [$H\cdots O$ = 2.51 (3) Å], $N-H\cdots N$ [$H\cdots N$ = 2.21 (3) Å] and $O-H\cdots N$ [$H\cdots O$ = 1.71 (3) Å]. In the crystal structure, molecules are linked into one-dimensional chains in the *a*-axis direction *via* intermolecular $O-H\cdots N$ [$H\cdots N$ = 1.95 (4) Å] hydrogen bonds.

Comment

There has been much interest in C_2 -symmetric N₂O₂ ligands and their metal complexes. Metal complexes based on the chiral ligand bis(salicylidene)-1,2,-diaminoethane, (1), have been used to catalyze a wide variety of organic reactions stereoselectively (Larrow & Jacobsen, 2004). Other C_2 symmetric N₂O₂ ligands with electron-withdrawing groups, (2), have recently been used as Ziegler-Natta catalysts in the synthesis of polyethylenes (Mason & Coates, 2004; Segal et al., 2005). However, compound (2) can, in principle, form three different isomers of octahedral complexes (*cis-\alpha*, *cis-\beta* and trans). More desirable are ligands which can form highly rigid single-isomer metal complexes. These ligands are particularly attractive for the development of catalysts and the investigation of stereoselective recognition of bidentate substrates. Here, we report the crystal structure of the title chiral C_2 symmetric N₂O₂ ligand, (3), which has two electron-withdrawing groups attached.



© 2005 International Union of Crystallography Printed in Great Britain – all rights reserved The connectivity of the two N and two O atoms in this tetradentate ligand is such that only the *cis*- α configuration of Received 28 October 2005 Accepted 17 November 2005 Online 26 November 2005

1958 reflections with $I > 2\sigma(I)$

 $R_{\rm int} = 0.070$

 $\theta_{\rm max} = 27.5^{\circ}$

 $h = -9 \rightarrow 10$

 $k = -11 \rightarrow 12$

 $l = -30 \rightarrow 30$



Figure 1

The structure of (3), showing 30% probability displacement ellipsoids. Dashed lines indicate hydrogen bonds.

the octahedral complex is possible upon deprotonation of the phenoxy group. A view of (3) is shown in Fig. 1 and selected torsion angles, describing the different orientations of the two dicyanoethyl groups, are given in Table 1.

In the molecule of (3), there are three intramolecular hydrogen bonds (Table 2), forming one S(5) and two S(6) rings (Bernstein *et al.*, 1995). In the crystal structure, molecules related by 2_1 screw axes are linked *via* O-H···N hydrogen bonds into one-dimensional C(10) chains in the *a*-axis direction (Fig. 2). These chains are further linked by weak C-H···O hydrogen bonds to form a three-dimensional network.

Experimental

1,2-(2-Hydroxyphenyl)-1,2-ethylenediamine was synthesized according to the literature procedure of Müller *et al.* (1989). Acrylonitrile (0.67 ml, 10.2 mmol) was added to a solution of (*S*,*S*)-1,2-(2-hydroxyphenyl)-1,2-ethylenediamine (1 g, 4.1 mmol) dissolved in methanol (14 ml). The mixture was stirred for 48 h at room temperature. After removal of the solvent, diethyl ether was added and the resulting solution was allowed to stand at room temperature for 3 d to give yellow crystals of (3). Spectroscopic analysis: ¹H NMR (400 MHz, CDCl₃, δ , p.p.m.): 2.50 (*m*, 4H), 2.84 (*m*, 4H), 3.99 (*s*, 2H), 6.74 (*t*, 2H), 6.81 (*d*, 2H), 6.87 (*d*, 2H), 7.18 (*t*, 2H).

Crystal data

Mo K α radiation Cell parameters from 16844 reflections $\theta = 2.6-27.5^{\circ}$ $\mu = 0.08 \text{ mm}^{-1}$ T = 150 (1) KPlate, colourless $0.22 \times 0.20 \times 0.02 \text{ mm}$





The crystal packing of (3), showing hydrogen bonds as dashed lines.

Data collection

Nonius KappaCCD diffractometer φ scans, and ω scans with κ offsets Absorption correction: multi-scan (*SORTAV*; Blessing, 1995) $T_{\min} = 0.723, T_{\max} = 0.999$ 16844 measured reflections 2492 independent reflections

Refinement

Refinement on F^2 $w = 1/[\sigma^2(F_0^2) + (0.05P)^2]$ $R[F^2 > 2\sigma(F^2)] = 0.039$ + 0.1038P] $wR(F^2) = 0.096$ where $P = (F_0^2 + 2F_c^2)/3$ S = 1.05 $(\Delta/\sigma)_{\rm max} = 0.001$ $\Delta \rho_{\rm max} = 0.14 \text{ e} \text{ Å}^{-3}$ 2492 reflections $\Delta \rho_{\rm min} = -0.17 \text{ e } \text{\AA}^{-3}$ 252 parameters H atoms treated by a mixture of Extinction correction: SHELXTL/ PC (Sheldrick, 2001) independent and constrained Extinction coefficient: 0.025 (3) refinement

Table 1

Selected torsion angles (°).

C1-N1-C2-C3	-179.39(17)	C11-N3-C12-C13	162.36 (18)
N1-C2-C3-C4	70.5 (2)	N3-C12-C13-C14	60.0 (3)
C5-C1-C11-N3	-178.44(16)		

Table 2			
Hydrogen-bond	geometry	(Å,	°)

$D-\mathrm{H}\cdots A$	D-H	$H \cdots A$	$D \cdots A$	$D - H \cdot \cdot \cdot A$
$01 - H10 \cdots N1$	1.00 (3)	1 71 (3)	2644(2)	154 (3)
$O2-H2O\cdots N4^{i}$	0.93 (4)	1.95 (4)	2.883 (3)	173 (3)
$N1 - H1N \cdot \cdot \cdot N3$	0.93 (3)	2.21 (3)	2.787 (3)	119 (2)
$N3 - H3N \cdots O2$	0.91 (3)	2.51 (3)	3.133 (3)	126 (2)
$C2-H2B\cdots N4^{ii}$	0.99	2.60	3.489 (3)	149
$C12-H12A\cdots O1^{iii}$	0.99	2.48	3.450 (3)	167
$C17-H17A\cdots N4^{i}$	0.95	2.56	3.308 (3)	135

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

H atoms bonded to N and O atoms were located in a difference map and refined isotropically. H atoms bonded to C atoms were placed in calculated positions, with C–H distances ranging from 0.95 to 1.00 Å. They were included in the refinement in the riding-model approximation, with $U_{iso}(H) = 1.2U_{eq}(C)$. In the absence of significant anomalous dispersion effects, Friedel pairs were merged before the refinement. The enantiomer was assigned from the known absolute configuration of the starting material.

Data collection: *COLLECT* (Nonius, 2003); cell refinement: *DENZO-SMN* (Otwinowski & Minor, 1997); data reduction: *DENZO-SMN*; program(s) used to solve structure: *SIR92* (Altomare *et al.*, 1994); program(s) used to refine structure: *SHELXTL/PC* (Sheldrick, 2001); molecular graphics: *SHELXTL/PC* and *PLATON* (Spek, 2003); software used to prepare material for publication: *SHELXTL/PC*.

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