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Key indicators

Single-crystal X-ray study T = 298 KMean σ (C–C) = 0.003 Å R factor = 0.043 wR factor = 0.110 Data-to-parameter ratio = 14.5

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

3-Chloro-N'-hydroxy-N-(1-naphthyl)benzenecarboximidamide

The molecule of the title compound, $C_{17}H_{13}ClN_2O$, adopts a *cis* configuration with respect to the C=N bond. The crystal structure features $O-H\cdots N$ hydrogen bonding.

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Comment

Amidoximes have been of great interest for many years. They have proved to be biologically active compounds in terms of antituberculostatic, antithrombotic and vasodilating, antimalarial, antidepressive and alpha-adrenergic activities (Nieves-Neira et al., 1999). They are also used as intermediates in the synthesis of various types of heterocyclic compounds, e.g. oxadiazoles (Yu et al., 2005) and oxadiazines (Dürüst, 1998). They are highly reactive compounds and have been extensively used as absorbents for uranium (Zhang et al., 2005), Cu²⁺ ions (Kondo et al., 1999) and vanadium (Seko et al., 2005) in sea-water. An amidoxime-based chelating resin has been prepared by Zohuriaan-Mehr et al. (2004) and exhibited high sorption capacity towards metal ions, e.g. Zn^{2+} , Cu^{2+} , Co^{2+} , Cd^{2+} and Ni^{2+} . As an extension of work on the synthesis of oxadiazoles, we report here the crystal structure of (I) (Fig. 1), a new amidoxime which is the Z(cis) isomer. In the crystal structure, the molecules are linked through weak O- $H \cdots N$ hydrogen bonds, forming a dimeric structure (Fig. 2).



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Figure 1

The molecular structure of (I), showing 50% probability displacement ellipsoids and the atom-numbering scheme.

Experimental

The reagents were purchased and used without further purification. 3-Chlorobenzohydroximinoyl chloride (6.01 g, 0.031 mol) was dissolved in dry CHCl₃ (50 ml), then a solution of 1-naphthylamine (9.04 g, 0.0632 mol) in CHCl₃ (10 ml) was added with stirring at room temperature. The reaction mixture was stirred for two days. The precipitate was filtered off and the excess solvent evaporated. The residue was subjected to flash column chromatography (hexane/ethyl acetate, 1:3), and allowed to evaporate slowly to give crystals.

Crystal data

$C_{17}H_{13}CIN_2O$
$M_r = 296.74$
Triclinic, P1
a = 7.7217 (5) Å
b = 8.7039 (4) Å
c = 11.9158 (4) Å
$\alpha = 104.099 \ (3)^{\circ}$
$\beta = 93.815 \ (4)^{\circ}$
$\gamma = 103.869 \ (4)^{\circ}$

Z = 2 $D_x = 1.319 \text{ Mg m}^{-3}$ Mo K α radiation $\mu = 0.26 \text{ mm}^{-1}$ T = 298 K Block, orange $0.52 \times 0.30 \times 0.20 \text{ mm}$

 $R_{\rm int} = 0.046$

 $\theta_{\rm max} = 25.5^{\circ}$

2764 independent reflections

1796 reflections with $I > 2\sigma(I)$

V = 747.41 (7) Å³

Data collection

Bruker–Nonius KappaCCD diffractometer φ and ω scans Absorption correction: none 10909 measured reflections

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_0^2) + (0.0431P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.043$	+ 0.1831P]
$wR(F^2) = 0.110$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.06	$(\Delta/\sigma)_{\rm max} = 0.001$
2764 reflections	$\Delta \rho_{\rm max} = 0.13 \ {\rm e} \ {\rm \AA}^{-3}$
190 parameters	$\Delta \rho_{\rm min} = -0.32 \ {\rm e} \ {\rm \AA}^{-3}$
H-atom parameters constrained	

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$O1-H1O\cdots N1^i$	0.88	1.98	2.768 (2)	147

Symmetry code: (i) -x + 2, -y, -z + 2.

All H atoms were located in a difference Fourier map. Aromatic H atoms were refined as riding on the respective C atoms with C-H = 0.93 Å and $U_{iso}(H) = 1.2U_{eq}(C)$. The O-H and N-H atoms were refined as riding in their as-found positions [O-H = 0.88 Å, N-H = 0.95 Å and $U_{iso}(H) = 1.2U_{eq}(N,O)$].

Data collection: COLLECT (Nonius, 1999); cell refinement: DIRAX/LSQ (Duisenberg, 1992); data reduction: EVALCCD



Figure 2

The centrosymmetric dimer of (I). Hydrogen bonds are indicated by dashed lines. H atoms not in involved in hydrogen bonds have been omitted.

(Duisenberg *et al.*, 2003); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *DIAMOND* (Brandenburg, 2006); software used to prepare material for publication: *publCIF* (Westrip, 2006).

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