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Key indicators

Single-crystal X-ray study T = 293 KMean σ (C–C) = 0.008 Å R factor = 0.056 wR factor = 0.128 Data-to-parameter ratio = 12.4

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

{Benzil bis[(2-oxidobenzylidene)hydrazone]}copper(II)

The title complex, $[Cu(C_{28}H_{20}N_4O_2)]$, can be described as a single-stranded helix with the Cu^{II} atom having a distorted square-planar coordination. Face-to-face and edge-to-face π - π stacking interactions and a C-H··· π interaction between the aromatic rings lead to a three-dimensional network in the crystal structure.

Comment

It is well established that self-assembly processes driven by metal-ligand interactions may generate a considerable variety of molecular architectures in a spontaneous and directed manner (Lehn, 1995; Su et al., 2003). Within the field of modern supramolecular chemistry, construction of coordination complexes via self-assembly of predesigned ligands with appropriate metal ions is a subject of current interest because of their unusual structures and special functional properties such as luminescence, redox activity and magnetism (Bai et al., 2004; Leininger et al., 2000). Bis-bidentate Schiff base ligands with two symmetrically related coordination capabilities derived from benzil dihydrazone are excellent ligands, which are easy to prepare and inexpensive, allowing us to probe systematically the effect of modifications to the ligand backbone through which we are attempting to control the precise topography, or microarchitecture, of the arrays (Bai et al., 2005; Chowdhury et al., 2003). In the present paper, a structural study of the title compound, (I), is presented.



The title complex, which contains one copper center and one ligand, forms a single-stranded helix (Fig. 1). The Cu^{II} atom has a highly distorted square-planar coordination, formed by two imine N atoms and two phenolate O atoms; angles at Cu are given in Table 1. The Cu–O and Cu–N distances agree well with those of related compounds (Fang *et al.*, 2001). The ligand is twisted, the C7–N1–N2–C8, C15–

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Figure 1

View of the molecular structure of (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 50% probability level.



Figure 2

Intermolecular face-to-face π - π stacking (dashed open lines), edge-toface $\pi - \pi$ stacking (dashed lines) and C – H··· π (dotted lines) interactions in (I), viewed along the b axis. H atoms, except that involved in the C- $H \cdot \cdot \pi$ interaction, have been omitted for clarity.

N3-N4-C22 and N2-C8-C15-N3 torsion angles being -122.7 (4), -134.9 (4) and -74.9 (6)°, respectively. The dihedral angle between the C9-C14 and C16-C21 phenyl rings is $88.9 (2)^{\circ}$. These results are similar to those found in a related compound (Chowdhury et al., 2003).

In the crystal structure of (I), the molecules further interact with each other, forming a three-dimensional network through face-to-face and edge-to-face π - π interactions and a C- $H \cdots \pi$ interaction (Fig. 2 and Table 2). The intermolecular face-to-face $\pi - \pi$ interaction between parallel salicylidenehydrazone groups, O1/C1-C7/N1/N2, related by an inversion center is characterized by a shortest contact distance of 3.281 (6) Å for C1···C7ⁱⁱ [symmetry code: (ii) 1 - x, 2 - y, (1 - z) and a centroid-centroid distance of 3.43 (6) Å. There are also extensive edge-to-face $\pi - \pi$ interactions between the benzene rings of adjacent molecules. The shortest contact distances are 3.482 (7) Å for C3···C23ⁱⁱⁱ [symmetry code: (iii) -1 + x, y, z], 3.594 (7) Å for C4...C10ⁱⁱⁱ and 3.581 (7) Å for $C11 \cdots C28^{iv}$ [symmetry code: (iv) x, 1 + y, z]. The dihedral angles between the two corresponding benzene rings are 55.3 (2)° for C1-C6 and C23ⁱⁱⁱ-C28ⁱⁱⁱ, 52.3 (2)° for C1-C6 and C9ⁱⁱⁱ–C14ⁱⁱⁱ and 82.2 (2)° for C9–C14 and C23^{iv}–C28^{iv}.

Experimental

All chemicals were of reagent grade quality obtained from commercial sources and used without further purification. Benzil bis(salicylidenehydrazone) (H_2L) was readily prepared by the reaction of benzil dihydrazone with salicylaldehyde in a 1:2 ratio. The ligand H₂L (0.2 mmol, 0.088 g) and Cu(ClO₄)₂·6H₂O (0.2 mmol, 0.074 g) were mixed in methanol (40 ml). The solution was left for 10 d at room temperature to afford green crystals (yield 80%). Analysis calculated for C₂₈H₂₀CuN₄O₂: C 66.2, H 4.0, N 11.0%; found: C 66.3, H 4.1, N 10.9%.

Crystal data

$[Cu(C_{28}H_{20}N_4O_2)]$	V = 1153.3 (4) Å ³
$M_r = 508.02$	Z = 2
Triclinic, P1	$D_x = 1.463 \text{ Mg m}^{-3}$
a = 9.223 (2) Å	Mo $K\alpha$ radiation
b = 10.702 (2) Å	$\mu = 0.98 \text{ mm}^{-1}$
c = 12.720 (3) Å	T = 293 (2) K
$\alpha = 85.132 \ (5)^{\circ}$	Block, green
$\beta = 89.749 \ (5)^{\circ}$	$0.20 \times 0.15 \times 0.15 \ \mathrm{mm}$
$\gamma = 67.273 \ (4)^{\circ}$	

Data collection

Bruker SMART CCD area-detector	5694 measured reflections
diffractometer	3934 independent reflections
φ and ω scans	2605 reflections with $I > 2\sigma(I)$
Absorption correction: multi-scan	$R_{\rm int} = 0.048$
(SADABS; Bruker, 2000)	$\theta_{\rm max} = 25.0^{\circ}$
$T_{\min} = 0.668, \ T_{\max} = 0.863$	

Refinement

Refinement on F^2	$\left[\exp\left\{2(\sin\theta/\lambda\right\}^2)\right]/[\sigma^2(F_0^2) +$
$R[F^2 > 2\sigma(F^2)] = 0.057$	$(0.045P)^2$]
$wR(F^2) = 0.128$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.02	$(\Delta/\sigma)_{\rm max} < 0.001$
3934 reflections	$\Delta \rho_{\rm max} = 0.64 \text{ e} \text{ \AA}^{-3}$
316 parameters	$\Delta \rho_{\rm min} = -0.27 \text{ e } \text{\AA}^{-3}$
H-atom parameters constrained	

Table 1

Selected geometric parameters (Å, $^\circ).$

Cu1—O1	1.891 (3)	Cu1-N1	1.939 (4)
Cu1—O2	1.876 (3)	Cu1-N4	1.976 (4)
O1-Cu1-O2	94.49 (15)	O2-Cu1-N1	152.98 (16)
O1-Cu1-N1	93.45 (15)	O2-Cu1-N4	92.74 (17)
O1-Cu1-N4	144.66 (14)	N1-Cu1-N4	95.59 (16)

Table 2Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
$C24-H24A\cdots Cg^{i}$	0.93	3.05	3.93 (2)	152
a				

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

All H atoms were positioned geometrically (C-H = 0.93 Å) and refined as riding, with $U_{iso}(H) = 1.2U_{eq}(C)$.

Data collection: *SMART* (Bruker, 2000); cell refinement: *SAINT* (Bruker, 2000); data reduction: *SAINT*; program(s) used to solve structure: *SHELXTL* (Bruker, 2000); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

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