

Bis{2'-[1-(2-hydroxy-5-chlorophenyl)ethylidene]-
benzohydrazido- κ^2O,N^2' }bis(pyridine- κN)zinc(II)Hapipah Ali,^{a*} Nur Ashikin
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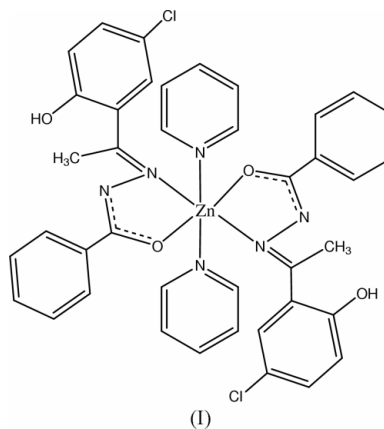
Key indicators

Single-crystal X-ray study
 $T = 273$ K
Mean $\sigma(C-C) = 0.005$ Å
 R factor = 0.054
 wR factor = 0.134
Data-to-parameter ratio = 16.1For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.

The title compound, $[Zn(C_{15}H_{13}ClN_2O_2)_2(C_5H_5N)_2]$, is a centrosymmetric monomer with octahedral geometry about the Zn atom. The molecule is stabilized by intermolecular C—H...O and C—H...Cl interactions, so forming a three-dimensional network.

Comment

Recrystallization of $[Zn(C_{13}H_9ClN_2O_2S)_2]$ from pyridine gave the dimeric compound $[Zn(C_{13}H_9ClN_2O_2S)_2(C_5H_5N)_2]$, with trigonal bipyramidal geometry at the central Zn atoms (Ali, Mohamad *et al.*, 2004). However, the title compound, $[Zn(C_{15}H_{13}ClN_2O_2)_2(C_5H_5N)_2]$, (I), which was also obtained by recrystallization of $[Zn(C_{15}H_{13}ClN_2O_2)_2]$ from pyridine, is monomeric with an octahedral geometry about the central Zn atom. The complex is also an analogue of the centrosymmetric compound $[Zn(C_{15}H_{13}N_2O_2)_2(C_5H_5N)_2]$ (Ali, Khamis *et al.*, 2004).



The molecular structure of (I) is illustrated in Fig. 1, and selected bond distances and angles are given in Table 1. The ligands are chelated in a bidentate manner *via* atoms O1 and N1, occupying the equatorial positions, the O1—Zn1—N1 bond angle being $103.47(8)^\circ$. The N atoms of the two pyridine molecules occupy the axial positions, the N3—Zn1—N3ⁱ bond angle being 180° [symmetry code: (i) $1 - x, -y, 1 - z$]. The structural dimensions of the ligand are in the normal range (Orpen *et al.*, 1979; Allen *et al.*, 1987) and in agreement with other octahedral zinc complexes (Tesouro Vallina & Stoeckli-Evans, 2002; Ali *et al.*, 2004). The chelated Zn1/O1/C7/N2/N1ⁱ fragment is planar, with a maximum deviation from the mean plane of $0.064(3)$ Å for atom N1ⁱ. The C8/C10—C15/O2/Cl1 fragment is planar [maximum deviation at C12 of $0.05(3)$ Å] and inclined to the C1—C7 fragment [maximum deviation $-0.011(4)$ Å for atom C5] by $43.53(15)^\circ$.

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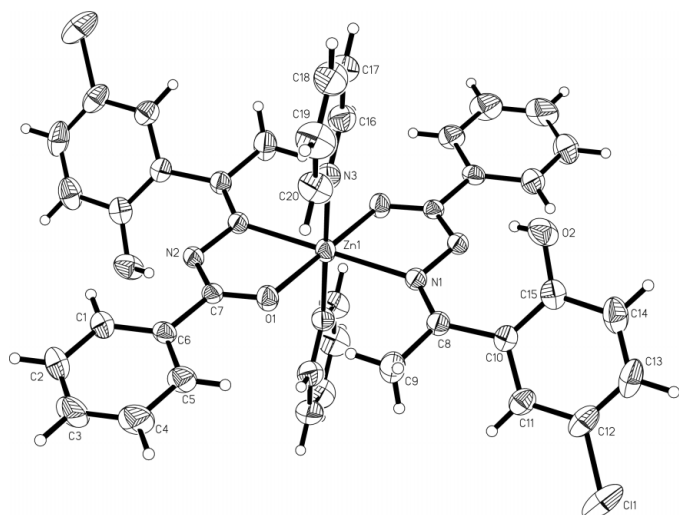


Figure 1
Molecular structure of (I), with 50% probability displacement ellipsoids. H atoms are represented by small spheres.

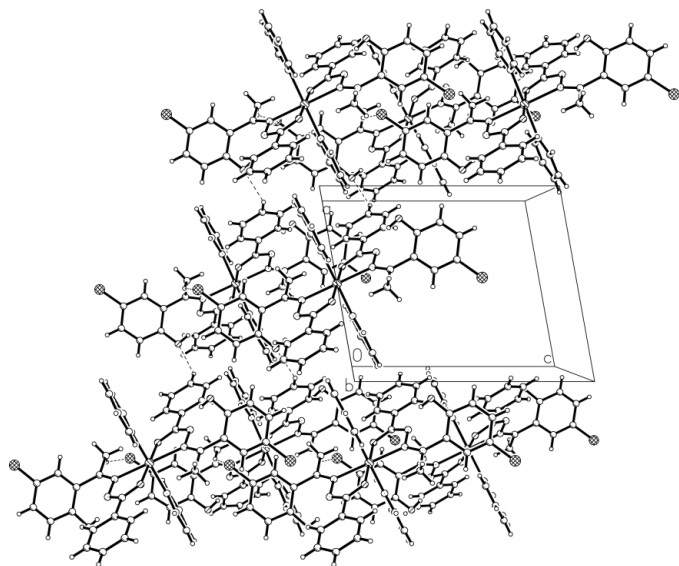


Figure 2
Packing diagram of the title complex, viewed down *b* axis. The dashed lines denote C—H...O and C—H...Cl interactions.

There are important intramolecular hydrogen-bonding interactions in the molecule, and details are given in Table 2. These are interactions of the types O—H...N, involving hydroxyl atom O2 and atom N2, and C—H...O, involving methyl group C9 and carbonyl atom O1 (O2—H2D...N2ⁱ and C9—H9A...O1). In the crystal, the molecules are linked by weak intermolecular C—H...O and C—H...Cl interactions (C3—H3A...O2ⁱⁱ and C9—H9B...Clⁱⁱⁱ), forming a three-dimensional network, as illustrated in Fig. 2.

Experimental

The complex was synthesized by the template condensation of 2-hydroxy-5-chloroacetophenonebenzhydrazide (0.3 g, 1.0 mmol) with zinc acetate dihydrate (0.11 g, 0.5 mmol), by refluxing and stirring in ethanol for 5 h. The light-yellow solid obtained was filtered off and recrystallized from pyridine.

Crystal data

[Zn(C₁₅H₁₃ClN₂O₂)₂(C₅H₅N)₂]
M_r = 799.00
 Monoclinic, *P*2₁/*c*
a = 10.2453 (17) Å
b = 15.388 (3) Å
c = 12.359 (2) Å
 β = 100.189 (3)°
V = 1917.8 (5) Å³
Z = 2

D_x = 1.384 Mg m⁻³
 Mo Kα radiation
 Cell parameters from 2652 reflections
 θ = 2.0–26.5°
 μ = 0.83 mm⁻¹
T = 273 (2) K
 Plate, light yellow
 0.48 × 0.46 × 0.09 mm

Data collection

Bruker SMART APEX area-detector diffractometer
 ω scans
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
T_{min} = 0.692, *T_{max}* = 0.929
 8429 measured reflections

3939 independent reflections
 3205 reflections with *I* > 2σ(*I*)
R_{int} = 0.021
 θ_{max} = 26.5°
h = -9 → 12
k = -13 → 19
l = -15 → 14

Refinement

Refinement on *F*²
R [*F*² > 2σ(*F*²)] = 0.054
wR (*F*²) = 0.134
S = 1.08
 3939 reflections
 245 parameters
 H atoms treated by a mixture of independent and constrained refinement

w = 1/[σ²(*F_o*²) + (0.0641*P*)² + 0.6133*P*]
 where *P* = (*F_o*² + 2*F_c*²)/3
 (Δ/σ)_{max} < 0.001
 Δρ_{max} = 0.54 e Å⁻³
 Δρ_{min} = -0.31 e Å⁻³

Table 1

Selected geometric parameters (Å, °).

Zn1—O1	2.022 (2)	N1—C8	1.291 (4)
Zn1—N1	2.206 (2)	N1—N2 ⁱ	1.393 (3)
Zn1—N3	2.243 (3)	N2—C7	1.321 (4)
Cl1—C12	1.737 (4)	N3—C16	1.324 (5)
O1—C7	1.263 (3)	N3—C20	1.331 (5)
O2—C15	1.348 (5)		
O1 ⁱ —Zn1—O1	180	O1—Zn1—N3	88.67 (10)
O1 ⁱ —Zn1—N1	76.53 (8)	N1—Zn1—N3	87.98 (10)
O1—Zn1—N1	103.47 (8)	N1 ⁱ —Zn1—N3	92.02 (10)
O1 ⁱ —Zn1—N3	91.33 (10)	N3—Zn1—N3 ⁱ	180

Symmetry code: (i) 1 - *x*, -*y*, 1 - *z*.

Table 2

Hydrogen-bonding geometry (Å, °).

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
C9—H9A...O1	0.96	2.24	3.177 (4)	165
O2—H2D...N2 ⁱ	0.87 (5)	1.69 (4)	2.522 (4)	160 (4)
C3—H3A...O2 ⁱⁱ	0.93	2.47	3.272 (6)	144
C9—H9B...Cl1 ⁱⁱⁱ	0.96	2.82	3.629 (5)	143

Symmetry codes: (i) 1 - *x*, -*y*, 1 - *z*; (ii) 1 + *x*, ½ - *y*, ½ + *z*; (iii) x, ½ - *y*, ½ + *z*.

All the H atoms could be located from Fourier difference maps. The hydroxyl H atom at O2 was refined isotropically. The remainder were included in calculated positions and treated as riding atoms [C—H_{ar} = 0.93 Å, with *U*_{iso}(H) = 1.2*U*_{eq}(C), and C—H₃ = 0.96 Å, with *U*_{iso}(H) = 1.5*U*_{eq}(C)].

Data collection: SMART (Siemens, 1996); cell refinement: SAINT (Siemens, 1996); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Sheldrick, 1997); software used to prepare material for

publication: *SHELXTL*, *PARST* (Nardelli, 1995) and *PLATON* (Spek, 2003).

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