

# Bis{ $\mu$ -*N*-[1-(2-oxidophenyl)ethylidene]benzohydrazido}bis[(dimethylsulfoxide)zinc(II)]

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## Key indicators

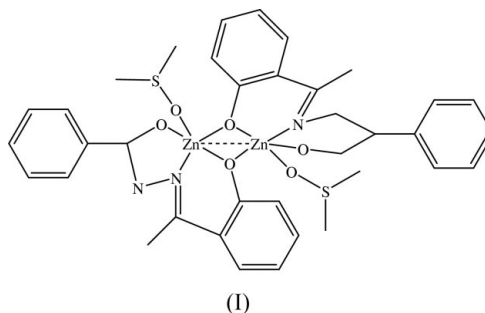
Single-crystal X-ray study  
T = 293 K  
Mean  $\sigma(C-C)$  = 0.002 Å  
R factor = 0.024  
wR factor = 0.068  
Data-to-parameter ratio = 17.2

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

The title compound,  $[Zn_2(C_{15}H_{12}N_2O_2)_2(C_2H_6OS)_2]$ , is a centrosymmetric dimer *via* a Zn—O bridge of 2.0038 (13) Å, with a Zn $\cdots$ Zn separation of 3.1370 (13) Å. The geometry of the five-coordinate environment of the Zn atoms is close to trigonal bipyramidal.

## Comment

It is known that recrystallization of some zinc complexes, such as Zn-3,5-diisopropylsalicylate in DMSO, has resulted in the formation of the DMSO derivative  $Zn(3,5-DIPS)_2(DMSO)_2$  (Morgant *et al.*, 1998). Similarly, the title compound, (I), was obtained when  $[Zn(C_{15}H_{12}N_2O_2)_2]$  was recrystallized from DMSO. However, unlike monomeric  $Zn(3,5-DIPS)_2(DMSO)_2$ , the crystal structure of (I) (Fig. 1) consists of centrosymmetric dimers, formed by linking two monomeric units through Zn1—O1<sup>i</sup> and Zn1<sup>i</sup>—O1 bonds of 2.0038 (13) Å (symmetry code in Table 1). The Zn1 $\cdots$ Zn1<sup>i</sup> separation is 3.1370 (13) Å. The relatively rare five-coordinated zinc complex shows that both central Zn atoms have geometries between square-pyramidal and trigonal bipyramidal, though they are closer to the latter.



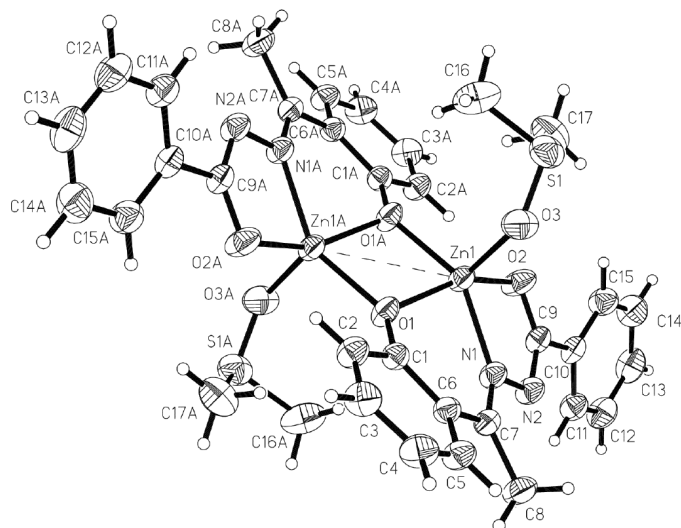
The bond distances and angles (Table 1) are normal (Orpen *et al.*, 1989; Allen *et al.*, 1987). The average Zn—O distance of 2.02 (2) Å is in agreement with the geometry of square pyramidal and trigonal bipyramidal, compared with 1.98 Å for tetrahedral (Morgant *et al.*, 1998) and 2.08 Å for octahedral (Babb *et al.*, 2003). The dimensions of the *O,N,O'*-tridentate ligand are typical of those of a Schiff base. The central Zn1/O1/Zn1<sup>i</sup>/O1<sup>i</sup> fragment is planar. The chelating O1/C1/C6/C7/N1 [maximum deviation is 0.045 (2) Å for C6] and O2/C9/N2/N1/C10 fragments are also planar, and inclined by 29.15 (8)° to each other. The two phenyl groups, C1—C6 and C9—C15, make a dihedral angle of 54.64 (8)°.

There is a weak C—H $\cdots$ O intramolecular contact (Table 2). In the crystal structure, the centrosymmetric dimers are linked by intermolecular interactions, C16—H16C $\cdots$ O3<sup>ii</sup>

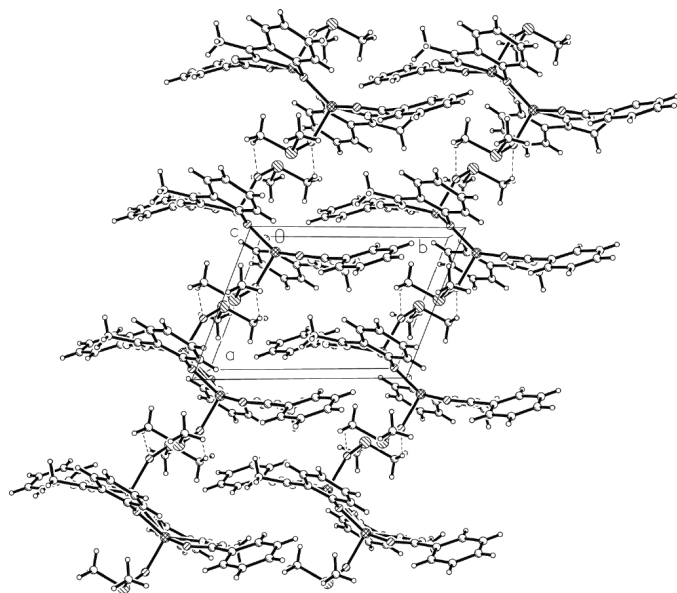
Received 3 July 2003

Accepted 21 July 2003

Online 31 July 2003



**Figure 1**  
The molecular structure of (I), shown with 50% probability displacement ellipsoids. The suffix *A* corresponds to symmetry code (i) in Tables 1 and 2.



**Figure 2**  
Packing diagram of the title complex, viewed down *c* axis. The dashed lines denote C—H...O intermolecular contacts.

(symmetry code as in Table 2) to form one-dimensional polymeric chains parallel to the *a* axis (Fig. 2).

## Experimental

The title complex was synthesized by the template condensation of benzhydrazide (2.00 g, 0.015 mol) and 2-hydroxyacetophenone (2.00 g, 0.015 mol) with zinc acetate dihydrate (1.65 g, 0.007 mol), with refluxing and stirring in ethanol for 5 h in the presence of triethylamine. The resulting pale-yellow solid was filtered and recrystallized from a minimum amount of DMSO. After standing at room temperature for 2 d, pale-yellow crystals were obtained.

## Crystal data

[Zn<sub>2</sub>(C<sub>15</sub>H<sub>12</sub>N<sub>2</sub>O<sub>2</sub>)<sub>2</sub>(C<sub>2</sub>H<sub>6</sub>OS)<sub>2</sub>]  
 $M_r = 791.53$   
 Triclinic,  $P\bar{1}$   
 $a = 7.690$  (4) Å  
 $b = 10.214$  (6) Å  
 $c = 11.643$  (6) Å  
 $\alpha = 98.479$  (13)°  
 $\beta = 94.262$  (12)°  
 $\gamma = 110.338$  (13)°  
 $V = 840.3$  (8) Å<sup>3</sup>

$Z = 1$   
 $D_x = 1.564$  Mg m<sup>-3</sup>  
 Mo  $K\alpha$  radiation  
 Cell parameters from 8764 reflections  
 $\theta = 1.7$ – $27.5$ °  
 $\mu = 1.60$  mm<sup>-1</sup>  
 $T = 293$  (2) K  
 Block, pale yellow  
 0.54 × 0.48 × 0.33 mm

## Data collection

Bruker SMART APEX CCD area-detector diffractometer  
 $\omega$  scans  
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)  
 $T_{\min} = 0.454$ ,  $T_{\max} = 0.588$   
 10556 measured reflections

3804 independent reflections  
 3653 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.019$   
 $\theta_{\max} = 27.5$ °  
 $h = -9 \rightarrow 9$   
 $k = -13 \rightarrow 13$   
 $l = -15 \rightarrow 15$

## Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.024$   
 $wR(F^2) = 0.068$   
 $S = 1.10$   
 3804 reflections  
 221 parameters  
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0374P)^2 + 0.2073P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} < 0.001$   
 $\Delta\rho_{\max} = 0.27$  e Å<sup>-3</sup>  
 $\Delta\rho_{\min} = -0.31$  e Å<sup>-3</sup>  
 Extinction correction: SHELXL97  
 Extinction coefficient: 0.0148 (15)

**Table 1**

Selected geometric parameters (Å, °).

Zn1—O1 <sup>i</sup>	2.0038 (13)	O2—C9	1.2834 (19)
Zn1—O2	2.0117 (14)	O3—S1	1.5142 (14)
Zn1—O3	2.0131 (16)	S1—C16	1.761 (2)
Zn1—N1	2.0466 (16)	S1—C17	1.781 (2)
Zn1—O1	2.0683 (15)	N1—C7	1.294 (2)
Zn1...Zn1 <sup>i</sup>	3.1370 (13)	N1—N2	1.3949 (17)
O1—C1	1.3391 (17)	N2—C9	1.309 (2)
O2—Zn1—O1	155.78 (5)	O2—Zn1—N1	78.70 (6)
O1 <sup>i</sup> —Zn1—O2	105.70 (6)	O3—Zn1—N1	107.52 (5)
O1 <sup>i</sup> —Zn1—O3	103.94 (6)	O1 <sup>i</sup> —Zn1—O1	79.24 (5)
O2—Zn1—O3	101.22 (6)	O3—Zn1—O1	100.46 (5)
O1 <sup>i</sup> —Zn1—N1	146.72 (5)	N1—Zn1—O1	84.59 (5)

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

**Table 2**

"C—H...O contacts (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>
C2—H2...O2 <sup>i</sup>	0.93	2.33	3.167 (3)	149
C16—H16C...O3 <sup>ii</sup>	0.96	2.42	3.258 (4)	145

Symmetry codes: (i)  $-x, -y, 1 - z$ ; (ii)  $1 - x, -y, 1 - z$ .

After their location in a difference map, all H atoms were placed geometrically at ideal positions and allowed to ride on the parent C atoms, with C—H = 0.93–0.96 Å and  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$  or  $1.5U_{\text{eq}}(\text{C})$ .

Data collection: SMART (Siemens, 1996); cell refinement: SAINT (Siemens, 1996); data reduction: SAINT; program(s) used to solve structure: SHELXTL (Sheldrick, 1997); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL, PARST (Nardelli, 1995) and PLATON (Spek, 2003).

The authors thank the Malaysian Government and both Universiti Malaya and Kebangsaan Malaysia for research grants IRPA Nos. 09-02-02-993.

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