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**Key indicators**

Single-crystal X-ray study  
*T* = 298 K  
 Mean  $\sigma$ (C–C) = 0.011 Å  
*R* factor = 0.074  
*wR* factor = 0.217  
 Data-to-parameter ratio = 14.2

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

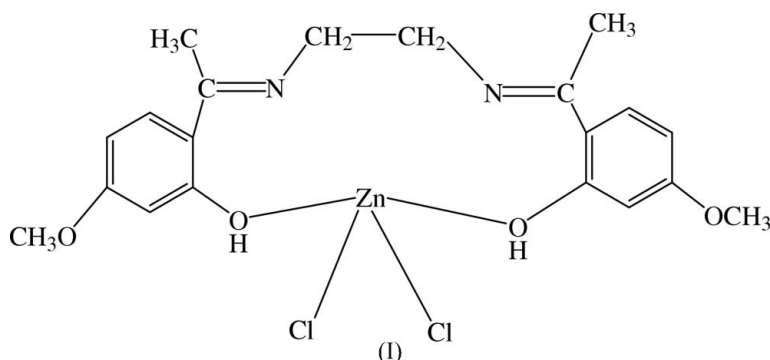
**{*N,N'*-Bis[1-(2-hydroxy-4-methoxyphenyl)ethylidene]ethylenediamine- $\kappa^2$ O,O'}**dichlorozinc(II)

In the title Schiff base complex, [ZnCl<sub>2</sub>(C<sub>10</sub>H<sub>12</sub>NO<sub>2</sub>)<sub>2</sub>], the Zn<sup>II</sup> ion is in a distorted tetrahedral ZnO<sub>2</sub>Cl<sub>2</sub> coordination environment.

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**Comment**

The Schiff base 2-hydroxy-4-methoxyacetophenone has potential biological properties. We report here the preparation and crystal structure of the title Zn<sup>II</sup> complex, (I), with 2-hydroxy-4-methoxyacetophenone.



The molecular structure of (I) is shown in Fig. 1. The Zn<sup>II</sup> ion is in a distorted tetrahedral coordination geometry (Table 1), formed by two hydroxyl O atoms and two Cl<sup>−</sup> anions. Atoms O1, O2 and N1 are almost coplanar with the C3–C8 benzene ring, while atoms O3, O4 and N2 are almost coplanar with the C12–C17 benzene ring. The dihedral angle between the two nine-atom mean planes is 14.6 (2)°. Intra-molecular hydrogen bonding occurs between the hydroxyl O atoms and imine N atoms (Table 2).

**Experimental**

To a stirred solution of 2-hydroxy-4-methoxyacetophenone (1.0 mmol) and ZnCl<sub>2</sub>·2H<sub>2</sub>O (1.0 mmol) in 30 ml absolute methanol was added dropwise a solution of ethylenediamine (1.0 mmol) in 10 ml absolute methanol at room temperature. After stirring for 2 h at 320 K, the precipitate was filtered off. Single crystals of (I) were obtained from the filtrate after 10 d.

*Crystal data*

[ZnCl <sub>2</sub> (C <sub>10</sub> H <sub>12</sub> NO <sub>2</sub> ) <sub>2</sub> ]	<i>Z</i> = 4
<i>M<sub>r</sub></i> = 492.68	<i>D<sub>x</sub></i> = 1.545 Mg m <sup>−3</sup>
Monoclinic, <i>P</i> 2 <sub>1</sub> / <i>c</i>	Mo <i>K</i> α radiation
<i>a</i> = 11.547 (3) Å	$\mu$ = 1.44 mm <sup>−1</sup>
<i>b</i> = 10.503 (3) Å	<i>T</i> = 298 (2) K
<i>c</i> = 17.469 (3) Å	Block, yellow
$\beta$ = 91.000 (3)°	0.22 × 0.14 × 0.10 mm
<i>V</i> = 2118.3 (9) Å <sup>3</sup>	

*Data collection*

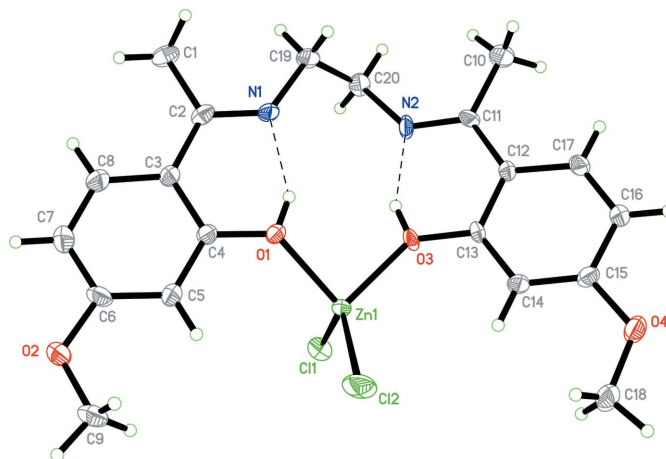
Bruker SMART CCD area-detector diffractometer  
 $\varphi$  and  $\omega$  scans  
 Absorption correction: multi-scan (SADABS; Sheldrick, 2002)  
 $T_{\min} = 0.742$ ,  $T_{\max} = 0.869$

10504 measured reflections  
 3729 independent reflections  
 2165 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.079$   
 $\theta_{\text{max}} = 25.0^\circ$

*Refinement*

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.074$   
 $wR(F^2) = 0.217$   
 $S = 1.03$   
 3729 reflections  
 262 parameters  
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.091P)^2 + 9.1479P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\text{max}} < 0.001$   
 $\Delta\rho_{\text{max}} = 0.88 \text{ e } \text{\AA}^{-3}$   
 $\Delta\rho_{\text{min}} = -0.87 \text{ e } \text{\AA}^{-3}$

**Figure 1**

The molecular structure of (I) with 30% probability displacement ellipsoids (arbitrary spheres for H atoms). Dashed lines indicate hydrogen bonds.

**Table 1**

Selected geometric parameters ( $\text{\AA}$ ,  $^\circ$ ).

Zn1—O1	1.975 (5)	Zn1—Cl1	2.233 (2)
Zn1—O3	1.957 (5)	Zn1—Cl2	2.219 (2)
O3—Zn1—O1	88.4 (2)	O3—Zn1—Cl1	108.8 (2)
O3—Zn1—Cl2	115.46 (18)	O1—Zn1—Cl1	108.74 (19)
O1—Zn1—Cl2	116.08 (19)	Cl2—Zn1—Cl1	116.02 (10)

**Table 2**

Hydrogen-bond geometry ( $\text{\AA}$ ,  $^\circ$ ).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
O1—H1 $\cdots$ N1	0.82	1.85	2.582 (8)	149
O3—H3 $\cdots$ N2	0.82	1.78	2.529 (8)	150

All H atoms were positioned geometrically and refined as riding on their parent atoms, with aromatic C—H distances of 0.93  $\text{\AA}$ , methyl C—H distances of 0.96  $\text{\AA}$ , methylene C—H distances of 0.97  $\text{\AA}$  and hydroxyl O—H distances of 0.82  $\text{\AA}$ , with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$  or  $1.5U_{\text{eq}}(\text{methyl C, O})$ .

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

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