

Mamiko Odoko,\* Natsuko  
Tsuchida and Nobuo OkabeFaculty of Pharmaceutical Sciences, Kinki  
University, Kowakae 3-4-1, Higashiosaka,  
Osaka 577-8502, JapanCorrespondence e-mail:  
odoko@phar.kindai.ac.jp

## Key indicators

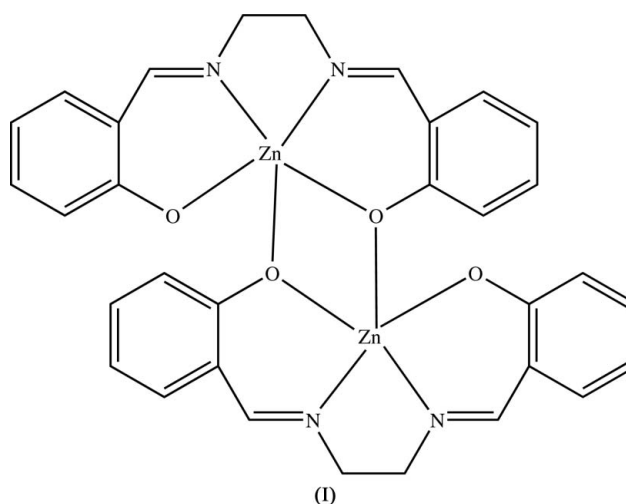
Single-crystal X-ray study  
 $T = 123$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.011$  Å  
 $R$  factor = 0.071  
 $wR$  factor = 0.200  
Data-to-parameter ratio = 16.0For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.Bis[ $\mu$ -2,2'-[ethane-1,3-diylbis(nitrilomethyl-  
idyne)]diphenolato}dizinc(II)

The crystal structure of the title compound,  $[\text{Zn}_2(\text{C}_{16}\text{H}_{14}\text{N}_2\text{O}_2)_2]$ , shows the dimeric complex to be centrosymmetric. Distorted pentacoordinate square-pyramidal geometry is formed around  $\text{Zn}^{\text{II}}$ , defined by two N atoms and two O atoms of the  $N,N'$ -bis(salicylidene)-1,2-ethylenediamine (salen) ligand and one bridging O atom of the adjacent ligand. Two salen ligands in this dimeric complex are twisted in opposite directions to each other.

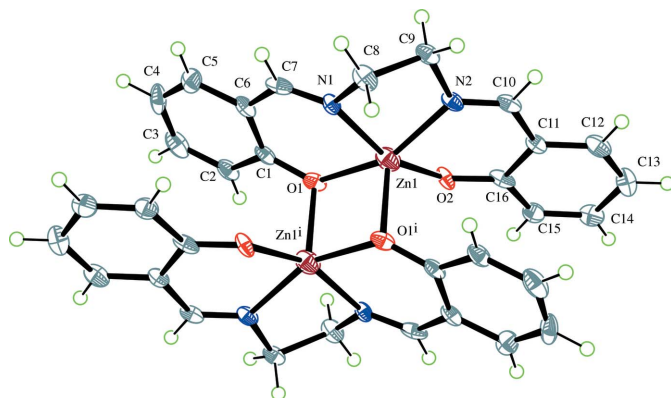
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## Comment

Zinc(II), a trace element for mammals, is essential for the various biochemical and physiological processes. Until now,  $\text{Zn}^{\text{II}}$  and its complexes have been found to exhibit *in vitro* insulinomimetic activity and anti-diabetic effects in animals (Chen *et al.*, 1998; Song *et al.*, 2001; Sakurai *et al.*, 2002). Several vanadium complexes with tetradentate Schiff base salen-type [salen is  $N,N'$ -bis(salicylidene)-1,2-ethylenediamine] ligands have been studied for use as insulin-enhancing agents (Durai & Saminathan, 1997; Correia *et al.*, 2004). In the present study, aimed to design new insulinomimetic  $\text{Zn}^{\text{II}}$  compounds, we have synthesized the title compound, (I), which has salen as the tetradentate Schiff base ligand, and analysed its crystal structure.



The crystal structure of (I) is shown in Fig. 1 with the atom-numbering scheme. Selected geometric parameters are given in Table 1. (I) is a centrosymmetric dimeric complex, rather than monomeric as in  $[\text{Zn}^{\text{II}}(\text{salen})\cdot\text{H}_2\text{O}]$  (Hall & Moore, 1966) and  $[\text{Zn}^{\text{II}}(\text{salen})\text{py}]$  (py = pyridine; Reglinski *et al.*, 2002).  $[\text{Zn}^{\text{II}}(\text{salen})]$  units in (I) are connected by the oxygen bridges  $\text{Zn1}-\text{O1}^i$  and  $\text{O1}-\text{Zn1}^i$  [symmetry code: (i)  $\frac{3}{2}-x, \frac{1}{2}-y, 1-z$ ].



**Figure 1**

A view of the structure of (I), showing the atom-numbering scheme. Displacement ellipsoids are shown at the 50% probability level and H atoms are shown as spheres of arbitrary radii. Symmetry code: (i)  $\frac{1}{2}-x$ ,  $\frac{1}{2}-y$ ,  $1-z$ .

z]. Distorted pentacoordinate square-pyramidal geometry is formed around  $\text{Zn}^{\text{II}}$ , defined by two N atoms and two O atoms of the salen ligand as the basal plane and one bridging O atom of the adjacent ligand as the apical atom.  $\text{Zn}^{\text{II}}$  sits in the body of the pyramid, 0.375 (3) Å from the O1–N1–N2–O2 basal plane. The two planar (r.m.s. deviation of fitted atoms = 0.0116 Å for O1/C1–C6/C7/N1 and 0.0425 Å for O10/C11–C16/C17/N2) Schiff base units are inclined at 21.9 (2)° in an umbrella-like conformation, similar to those in  $[\text{Zn}^{\text{II}}(\text{salen})\cdot\text{H}_2\text{O}]$  and  $[\text{Zn}^{\text{II}}(\text{salen})\text{py}]$ . The two salen ligands in (I) are twisted in opposite directions to each other. The ethylenediamine group of the salen ligand is in a *gauche* conformation  $[\text{N1}–\text{C8}–\text{C9}–\text{N2} = 45.0 (7)^\circ]$ , as seen in  $[\text{Zn}^{\text{II}}(\text{salen})\cdot\text{H}_2\text{O}]$ .

The crystal packing of (I) is stabilized by van der Waals interactions between neighboring complexes.

## Experimental

*N,N'*-Bis(salicylidene)-1,2-ethylenediamine (5 mg) was dissolved in MeOH (5 ml) and zinc perchlorate hexahydrate (3.4 mg) dissolved in a small amount of water was added at room temperature. Colorless needle-shaped crystals appeared from this mixture after evaporation over a few days at room temperature.

### Crystal data

$[\text{Zn}_2(\text{C}_{16}\text{H}_{14}\text{N}_2\text{O}_2)_2]$	$D_x = 1.646 \text{ Mg m}^{-3}$
$M_r = 663.36$	Mo $K\alpha$ radiation
Monoclinic, $C2/c$	Cell parameters from 6984 reflections
$a = 27.06 (5) \text{ \AA}$	$\theta = 3.0\text{--}27.5^\circ$
$b = 6.79 (1) \text{ \AA}$	$\mu = 1.84 \text{ mm}^{-1}$
$c = 14.70 (3) \text{ \AA}$	$T = 123.1 \text{ K}$
$\beta = 97.67 (7)^\circ$	Needle, colorless
$V = 2677 (8) \text{ \AA}^3$	$0.60 \times 0.05 \times 0.05 \text{ mm}$
$Z = 4$	

### Data collection

Rigaku R-AXIS RAPID diffractometer	3063 independent reflections
$\omega$ scans	919 reflections with $F^2 > 2\sigma(F^2)$
Absorption correction: multi-scan (ABSCOR; Higashi, 1995)	$R_{\text{int}} = 0.158$
$T_{\text{min}} = 0.197$ , $T_{\text{max}} = 0.912$	$\theta_{\text{max}} = 27.5^\circ$
12653 measured reflections	$h = -34 \rightarrow 34$
	$k = -8 \rightarrow 8$
	$l = -19 \rightarrow 19$

### Refinement

Refinement on $F^2$	All H-atom parameters refined
$R[F^2 > 2\sigma(F^2)] = 0.071$	$w = 1/[\sigma^2(F_o^2) + (0.0929P)^2]$
$wR(F^2) = 0.200$	where $P = (F_o^2 + 2F_c^2)/3$
$S = 0.89$	$(\Delta/\sigma)_{\text{max}} = 0.001$
3063 reflections	$\Delta\rho_{\text{max}} = 0.97 \text{ e \AA}^{-3}$
191 parameters	$\Delta\rho_{\text{min}} = -1.06 \text{ e \AA}^{-3}$

**Table 1**

Selected geometric parameters (Å, °).

Zn1–O1	2.053 (5)	Zn1–N1	2.072 (5)
Zn1–O1 <sup>i</sup>	2.053 (5)	Zn1–N2	2.078 (6)
Zn1–O2	1.973 (4)		
O1–Zn1–O1 <sup>i</sup>	88.3 (2)	O1 <sup>i</sup> –Zn1–N1	102.2 (2)
O1–Zn1–O2	98.0 (2)	O1 <sup>i</sup> –Zn1–N2	113.0 (2)
O1–Zn1–N1	86.2 (2)	O2–Zn1–N2	87.4 (2)
O1 <sup>i</sup> –Zn1–O2	100.4 (2)	N1–Zn1–N2	80.5 (2)

Symmetry code: (i)  $-x + \frac{3}{2}$ ,  $-y + \frac{1}{2}$ ,  $-z + 1$ .

All H atoms were located in difference Fourier maps and then were placed in ideal positions and refined as riding [ $\text{C}–\text{H} = 0.93 \text{ \AA}$  and  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$  for aromatic H atoms;  $\text{C}–\text{H} = 0.97 \text{ \AA}$  and  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$  for methine and methylene H atoms]. The crystal used for data collection was very poorly diffracting but it was the best that could be found. The highest electron-density peak was located close to atoms Zn1 and O2.

Data collection: *RAPID-AUTO* (Rigaku, 1998); cell refinement: *RAPID-AUTO*; data reduction: *CrystalStructure* (Rigaku/MS, 2005) and *CRYSTALS* (Betteridge *et al.*, 2003); program(s) used to solve structure: *SIR97* (Altomare *et al.*, 1999); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3* (Farrugia, 1997); software used to prepare material for publication: *CrystalStructure*.

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