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## Key indicators

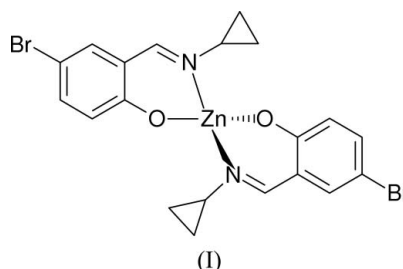
Single-crystal X-ray study  
 $T = 298$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.007$  Å  
 $R$  factor = 0.043  
 $wR$  factor = 0.102  
Data-to-parameter ratio = 18.9For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.Bis[4-bromo-2-(cyclopropyliminomethyl)-  
phenolato]zinc(II)In the mononuclear title complex,  $[\text{Zn}(\text{C}_{10}\text{H}_9\text{BrNO})_2]$ , the  $\text{Zn}^{\text{II}}$  atom is four-coordinated by two imine N and two phenolate O atoms from two Schiff base ligands in a distorted tetrahedral geometry.

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## Comment

As an extension of the work on the structural characterization of Schiff base zinc(II) compounds (You, 2005*a,b,c*), the title Schiff base zinc(II) complex, (I), is reported.

Complex (I) is a mononuclear zinc(II) compound (Fig. 1). The bond lengths and angles (Table 1) are comparable to those observed in bis[2-(cyclopropyliminomethyl)phenolato]zinc(II) (You *et al.*, 2003). The  $\text{Zn}^{\text{II}}$  atom is coordinated by two N and two O atoms from two Schiff base ligands. This  $\text{ZnN}_2\text{O}_2$  coordination forms a distorted tetrahedral geometry, with angles subtended at the  $\text{Zn}^{\text{II}}$  atom in the range 96.69 (12)–120.01 (14)°. C—H...O intermolecular hydrogen bonds (Table 2) link symmetry-related molecules into a three-dimensional network.

## Experimental

Cyclopropylamine (0.1 mmol, 5.7 mg) and 5-bromosalicylaldehyde (0.1 mmol, 20.1 mg) were dissolved in MeOH (10 ml). The mixture was stirred at room temperature for 10 min to give a clear yellow solution. To this solution was added a MeOH solution (5 ml) of  $\text{Zn}(\text{CH}_3\text{COO})_2 \cdot 4\text{H}_2\text{O}$  (0.1 mmol, 25.6 mg), with stirring. The resulting mixture was stirred for another 10 min at room temperature. After keeping the filtrate in air for 7 d, colourless block-shaped crystals were formed at the bottom of the vessel.

## Crystal data

$[\text{Zn}(\text{C}_{10}\text{H}_9\text{BrNO})_2]$   
 $M_r = 543.55$   
Orthorhombic,  $Pbca$   
 $a = 23.260$  (2) Å  
 $b = 13.305$  (2) Å  
 $c = 13.119$  (3) Å  
 $V = 4060.0$  (12) Å<sup>3</sup>  
 $Z = 8$   
 $D_x = 1.779$  Mg m<sup>-3</sup>

Mo  $K\alpha$  radiation  
Cell parameters from 4849  
reflections  
 $\theta = 2.3$ – $23.6^\circ$   
 $\mu = 5.16$  mm<sup>-1</sup>  
 $T = 298$  (2) K  
Block, colourless  
0.24 × 0.20 × 0.12 mm

Data collection

Bruker SMART CCD area-detector diffractometer  
 $\omega$  scans  
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)  
 $T_{\min} = 0.346$ ,  $T_{\max} = 0.538$   
 21839 measured reflections

4619 independent reflections  
 2693 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.065$   
 $\theta_{\text{max}} = 27.5^\circ$   
 $h = -29 \rightarrow 28$   
 $k = -16 \rightarrow 17$   
 $l = -16 \rightarrow 16$

Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.043$   
 $wR(F^2) = 0.102$   
 $S = 1.01$   
 4619 reflections  
 244 parameters  
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0401P)^2 + 2.1967P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\text{max}} = 0.001$   
 $\Delta\rho_{\text{max}} = 0.55 \text{ e } \text{\AA}^{-3}$   
 $\Delta\rho_{\text{min}} = -0.78 \text{ e } \text{\AA}^{-3}$

Table 1

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

Zn1—O2	1.904 (3)	Zn1—N1	2.002 (3)
Zn1—O1	1.916 (3)	Zn1—N2	2.013 (3)
O2—Zn1—O1	113.95 (12)	O2—Zn1—N2	97.50 (12)
O2—Zn1—N1	116.46 (13)	O1—Zn1—N2	113.33 (13)
O1—Zn1—N1	96.69 (12)	N1—Zn1—N2	120.01 (14)

Table 2

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
C8—H8 $\cdots$ O1 <sup>i</sup>	0.98	2.54	3.488 (5)	162
C18—H18 $\cdots$ O1 <sup>ii</sup>	0.98	2.49	3.442 (5)	163
C20—H20B $\cdots$ O1	0.97	2.55	3.419 (7)	149

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

H atoms were placed in idealized positions and constrained to ride on their parent atoms, with  $C-H = 0.93-0.98 \text{ \AA}$  and  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ .

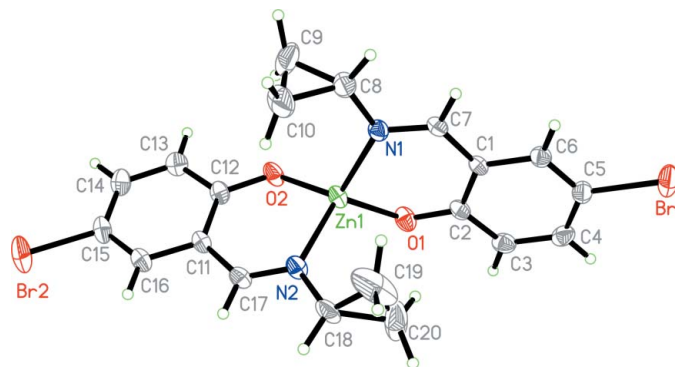


Figure 1

The structure of (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level.

Data collection: SMART (Bruker, 1998); cell refinement: SAINT (Bruker, 1998); 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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