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Key indicators

Single-crystal X-ray study

T = 273 K

Mean $\sigma(\text{C}-\text{C}) = 0.007 \text{ \AA}$

R factor = 0.055

wR factor = 0.130

Data-to-parameter ratio = 18.5

For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.Bis{2-[(2-aminophenyl)iminomethyl]-
6-carboxyphenolato- $\kappa^3\text{O},\text{N},\text{N}'$ }zinc(II)

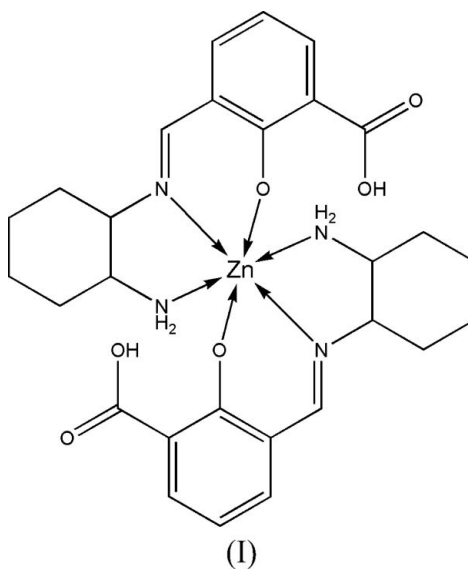
The title compound, $[\text{Zn}(\text{C}_{14}\text{H}_{17}\text{N}_2\text{O}_3)_2]$, is a complex of Zn^{II} and the Schiff base 2-[(2-aminophenyl)iminomethyl]-6-carboxyphenol synthesized by the reaction of 3-carboxylsalicylaldehyde and (1*R*,2*R*)-diaminocyclohexane in equimolar ratio. In the crystal structure, the Zn^{II} ion is six-coordinated in an octahedral environment (N_4O_2) formed by two monovalent anions. Each anion acts as a tridentate *NON*-chelator and occupies a coordination plane. The two *NON* coordination planes formed are nearly perpendicular to each other.

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Comment

Complexes of Schiff bases play an important role in modern chemistry, because they show abundant bioactivity (Dios *et al.*, 2002), medicinal activity (Sharma, 2004) and catalytic activity. The Schiff bases of 3-carboxylsalicylaldehyde and diamines (1,2-diaminoethane, 1,2-diaminopropane, 1,3-diaminopropane, *etc.*) have many coordination modes, so they can form many mono- and polynuclear complexes (Tao *et al.*, 2002; Li *et al.*, 2006). Recently, our interest has been focused on asymmetric Schiff bases and their complexes (Zhang *et al.*, 2005; Yu *et al.*, 2006). Here, we report the crystal structure of the title mononuclear zinc(II) complex, (I), in which the ligand is the Schiff base of (1*R*,2*R*)-diaminocyclohexane monocondensed with one equivalent of 3-carboxylsalicylaldehyde.



Complex (I) is formed by a Zn^{II} cation and two univalent anions. The Zn^{II} cation is coordinated by four N atoms and two O atoms in an octahedral geometry. Each anionic ligand

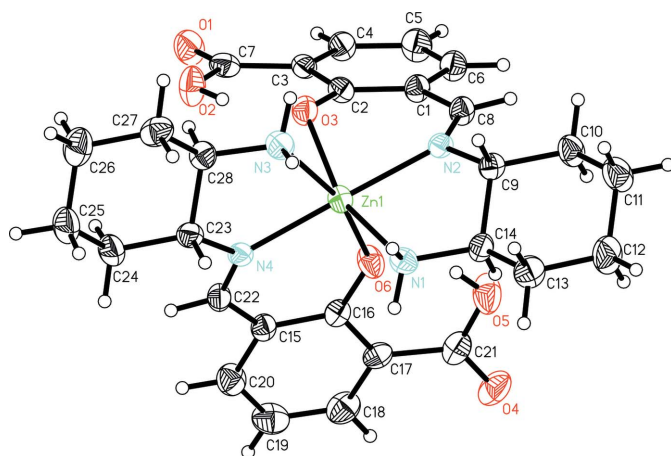


Figure 1
The molecular structure of (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 30% probability level.

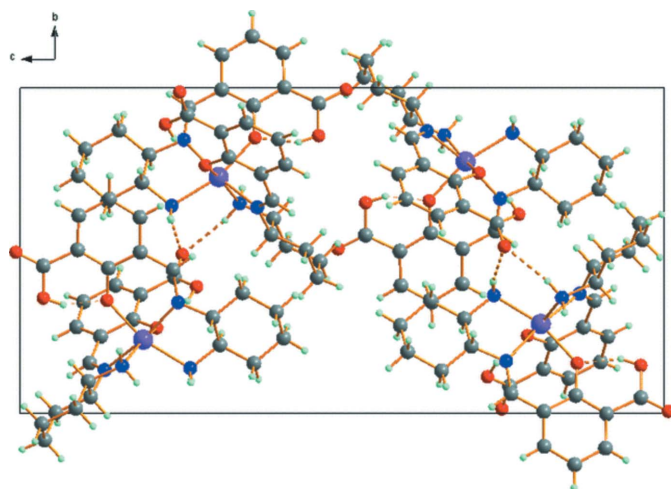


Figure 2
The molecular packing of (I), viewed along the *a* axis. Hydrogen bonds are indicated by dashed lines.

acts as a tridentate chelator *via* two N atoms and one phenolate O atom, and occupies a coordination plane. The two NON coordination planes formed are nearly perpendicular to each other, with a dihedral angle of 88.08 (13)°.

The carboxy group in the ligand does not coordinate to the central atom, but forms an intramolecular hydrogen bond with the phenolate O atom. Intermolecular hydrogen bonds are formed through the H atoms of the amine groups to carboxyl O atoms and link the molecules into chains running parallel to the *b* axis (Table 1, Fig. 2).

Experimental

3-Carboxylsalicylaldehyde (0.332 g, 2.0 mmol) was dissolved in methanol (30 ml). To this solution, an aqueous solution (10 ml) containing NaOH (0.160 g, 4.0 mmol) and cyclohexane-1,2-diammonium (1*R*,2*R*)-tartrate (0.5286 g, 2.0 mmol) was added. The mixed solution was stirred at 333 K for 1 h and then filtered to give a clear yellow solution. To this yellow solution, zinc acetate [$\text{Zn}(\text{OOCCH}_3)_2 \cdot 2\text{H}_2\text{O}$; 0.439 g, 2.0 mmol] dissolved in methanol

(30 ml) was added. The system was stirred at 333 K for 2 h, then cooled and filtered. The filtrate was sealed and kept at room temperature. After 30 d, yellow block-shaped crystals of (I), suitable for X-ray diffraction, were obtained.

Crystal data

$[\text{Zn}(\text{C}_{14}\text{H}_{17}\text{N}_2\text{O}_3)_2]$	$V = 2793.5 (11) \text{ \AA}^3$
$M_r = 587.96$	$Z = 4$
Orthorhombic, $P2_12_12_1$	Mo $K\alpha$ radiation
$a = 8.858 (2) \text{ \AA}$	$\mu = 0.93 \text{ mm}^{-1}$
$b = 12.610 (3) \text{ \AA}$	$T = 273 (2) \text{ K}$
$c = 25.008 (6) \text{ \AA}$	$0.15 \times 0.12 \times 0.11 \text{ mm}$

Data collection

Bruker APEXII CCD area-detector diffractometer	17440 measured reflections
Absorption correction: multi-scan (SADABS; Bruker, 2005)	6561 independent reflections
$T_{\min} = 0.873$, $T_{\max} = 0.905$	3447 reflections with $I > 2\sigma(I)$
	$R_{\text{int}} = 0.083$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.055$	$\Delta\rho_{\max} = 0.52 \text{ e \AA}^{-3}$
$wR(F^2) = 0.130$	$\Delta\rho_{\min} = -0.50 \text{ e \AA}^{-3}$
$S = 0.97$	Absolute structure: Flack (1983), with 3755 Friedel pairs
6561 reflections	Flack parameter: 0.018 (17)
354 parameters	
H-atom parameters constrained	

Table 1

Hydrogen-bond geometry (\AA , °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{O5}-\text{H5}\cdots\text{O6}$	0.82	1.71	2.473 (4)	153
$\text{O2}-\text{H2}\cdots\text{O3}$	0.82	1.67	2.439 (5)	155
$\text{N3}-\text{H3A}\cdots\text{O1}^i$	0.90	2.37	3.253 (6)	168
$\text{N1}-\text{H1A}\cdots\text{O1}^i$	0.90	2.22	3.114 (5)	170

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

All H atoms were included in calculated positions, with C–H distances ranging from 0.93 to 0.98 \AA , N–H distances of 0.90 \AA and O–H distances of 0.82 \AA . They were refined in the riding-model approximation, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C}, \text{N})$ or $1.5U_{\text{eq}}(\text{O})$.

Data collection: APEX2 (Bruker, 2005); cell refinement: SAINT (Bruker, 2005); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Bruker, 2005); software used to prepare material for publication: SHELXTL.

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