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Key indicators

Single-crystal X-ray study
T = 293 K
Mean $\sigma(\text{C}-\text{C}) = 0.004 \text{ \AA}$
R factor = 0.036
wR factor = 0.089
Data-to-parameter ratio = 16.9

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

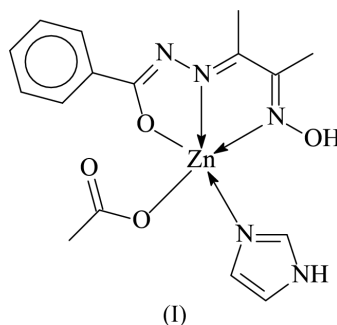
(Acetato- κO)[N-(diacetyl monooxime)-
N'-benzoylhydrazonato- $\kappa^3\text{O},\text{N},\text{N}'$]-
(1H-imidazole- κN^3)zinc(II)

In the title compound, $[\text{Zn}(\text{C}_2\text{H}_3\text{O}_2)(\text{C}_{11}\text{H}_{12}\text{N}_3\text{O}_2)(\text{C}_3\text{H}_4\text{N}_2)]$, the Zn atom is five-coordinate in a ZnN_3O_2 trigonal-bipyramidal environment. Adjacent molecules are linked by an intermolecular hydrogen bond into a chain running along the *a* axis of the monoclinic cell.

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Comment

A previous study has documented the zinc complex of the hydrazone that was synthesized by condensing benzoylacetone with 4-methoxybenzoylhydrazine; the five-coordinate metal atom shows *trans*-trigonal geometry (Gao *et al.*, 2004). The Zn complex of N-(diacetylmonooxime)-N'-(benzoyl)-hydrazone crystallizes with imidazole, but the complex, (I), has the metal atom in a geometry that is intermediate between a square pyramid and a trigonal bipyramid (Fig. 1). The bond angles at the Zn atom of (I) range from 72.65 (7) to 148.61 (6)°.



The ligand has been characterized previously in a Cu complex (Chumakov *et al.*, 1979). The bond lengths of the

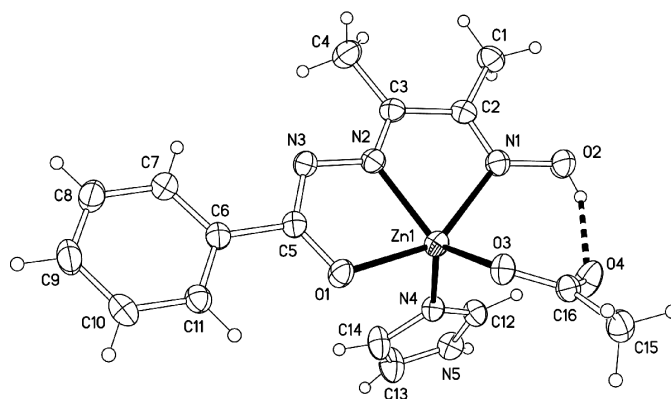


Figure 1
A view of (I), showing the atom-numbering scheme and with 30% probability displacement ellipsoids. H atoms are shown as small spheres of arbitrary radii. The dashed line indicates the intramolecular hydrogen bond.

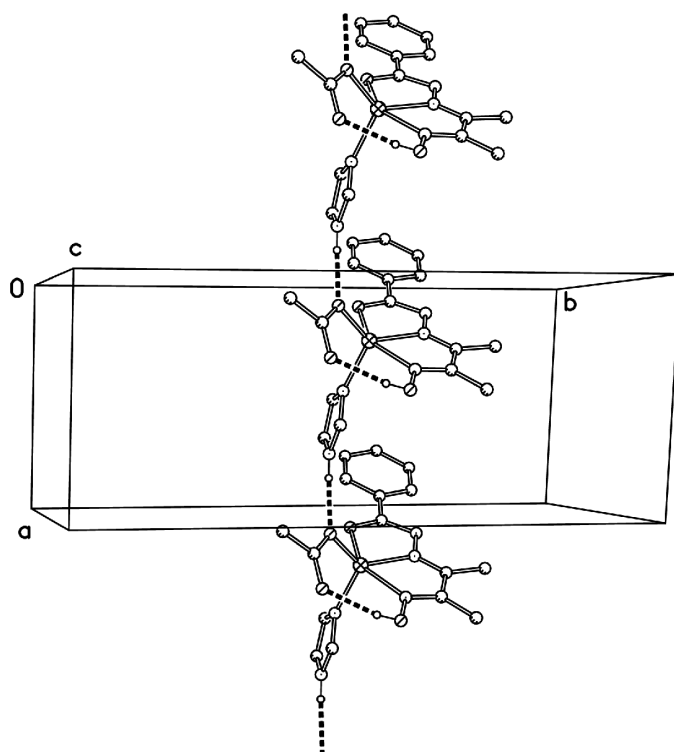


Figure 2
A view of the hydrogen-bonded chain structure of (I).

hydrazone moiety in (I) suggest delocalization of electrons throughout. The oxime atom O2 of the hydrazone ligand forms an intramolecular O—H...O hydrogen bond with the uncoordinated acetate O atom. Another hydrogen bond gives rise to the formation of a linear chain along the *a* direction. (Table 2, Fig. 2).

Experimental

The hydrazone ligand was synthesized by condensing benzoylhydrazine with an equimolar quantity of diacetyl monooxime in ethanol. A methanol solution containing zinc(II) acetate dihydrate (1 mmol) and imidazole (1 mmol) was added to a methanol solution (25 ml) of diacetylmonooxime benzoylhydrazine (1 mmol). The resulting mixture was refluxed with stirring for 30 min, cooled slowly to room temperature and then filtered. Yellow crystals of (I) were isolated from the solution after several days. Analysis calculated for $C_{16}H_{19}N_5O_4Zn$: C 46.79, H 4.66, N 17.05%; found: C 47.52, H 4.83, N 16.92%.

Crystal data

$[Zn(C_2H_3O_2)(C_{11}H_{12}N_5O_2) \cdot (C_3H_4N_2)]$
 $M_r = 410.75$
 Monoclinic, $P2_1/c$
 $a = 7.731(2) \text{ \AA}$
 $b = 18.334(4) \text{ \AA}$
 $c = 12.931(3) \text{ \AA}$
 $\beta = 100.07(3)^\circ$
 $V = 1804.6(8) \text{ \AA}^3$
 $Z = 4$

$D_x = 1.512 \text{ Mg m}^{-3}$
 Mo $K\alpha$ radiation
 Cell parameters from 11 353 reflections
 $\theta = 3.4\text{--}27.4^\circ$
 $\mu = 1.39 \text{ mm}^{-1}$
 $T = 293(2) \text{ K}$
 Prism, yellow
 $0.37 \times 0.25 \times 0.18 \text{ mm}$

Data collection

Rigaku R-Axis RAPID diffractometer	4122 independent reflections
ω scans	3431 reflections with $I > 2\sigma(I)$
Absorption correction: multi-scan (ABSCOR; Higashi, 1995)	$R_{\text{int}} = 0.033$
$T_{\text{min}} = 0.626$, $T_{\text{max}} = 0.788$	$\theta_{\text{max}} = 27.5^\circ$
16 947 measured reflections	$h = -10 \rightarrow 10$
	$k = -23 \rightarrow 23$
	$l = -16 \rightarrow 16$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.036$
 $wR(F^2) = 0.089$
 $S = 1.04$
 4122 reflections
 244 parameters
 H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0509P)^2 + 0.4702P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.47 \text{ e \AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.26 \text{ e \AA}^{-3}$

Table 1

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

Zn1—N1	2.272 (2)	N2—N3	1.374 (2)
Zn1—N2	2.037 (2)	N2—C3	1.290 (3)
Zn1—N4	2.000 (2)	N3—C5	1.329 (3)
Zn1—O1	2.100 (2)	O1—C5	1.274 (2)
Zn1—O3	1.972 (1)	O2—N1	1.382 (2)
N1—C2	1.279 (3)	C2—C3	1.485 (3)
N2—Zn1—N1	72.65 (7)	O1—Zn1—N1	148.61 (6)
N2—Zn1—O1	76.23 (6)	O3—Zn1—N1	101.48 (7)
N4—Zn1—N1	96.71 (7)	O3—Zn1—N2	130.82 (6)
N4—Zn1—N2	112.70 (7)	O3—Zn1—N4	116.48 (6)
N4—Zn1—O1	98.95 (7)	O3—Zn1—O1	95.60 (6)

Table 2

Hydrogen-bonding geometry (\AA , $^\circ$).

$D\text{—}H\cdots A$	$D\text{—}H$	$H\cdots A$	$D\cdots A$	$D\text{—}H\cdots A$
N5—H18...O3 ⁱ	0.85 (3)	1.91 (3)	2.756 (2)	176 (3)
O2—H17...O4	0.84 (3)	1.99 (3)	2.789 (2)	159 (4)

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

The H atoms on C atoms were placed in calculated positions, with C—H = 0.93 (aromatic) or 0.96 \AA (methyl), and were refined using a riding model, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{aromatic C})$ or $1.5U_{\text{eq}}(\text{methyl C})$. The torsion angle of the methyl groups was refined. The H atoms on the imidazole N and oxime O atoms were located in a difference Fourier map and refined with N—H and O—H distances restrained to 0.86 (1) and 0.85 (1) \AA , respectively, and with $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{N,O})$.

Data collection: *RAPID-AUTO* (Rigaku, 1998); cell refinement: *RAPID-AUTO*; data reduction: *CrystalStructure* (Rigaku/MS, 2002); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEPII* (Johnson, 1976); software used to prepare material for publication: *SHELXL97*.

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