

Diaquabis[5-(pyrazin-2-yl- κN^1)-3-(pyridin-3-yl)-1,2,4-triazolido- κN^1]-zinc

Ye-Nan Wang and Wen-Wen Dong*

College of Materials & Chemical Engineering, China Three Gorges University, Yichang 443002, People's Republic of China
Correspondence e-mail: dongww1@126.com

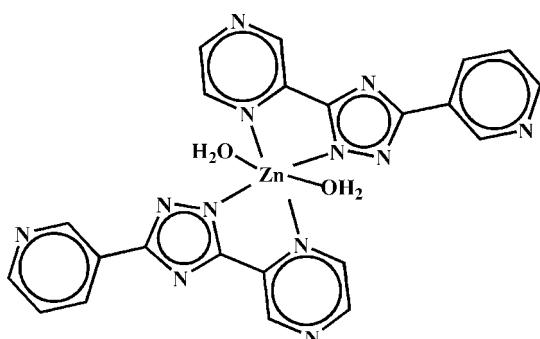
Received 20 February 2014; accepted 23 February 2014

Key indicators: single-crystal X-ray study; $T = 296\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$; R factor = 0.037; wR factor = 0.110; data-to-parameter ratio = 14.2.

In the title compound, $[\text{Zn}(\text{C}_{11}\text{H}_7\text{N}_6)_2(\text{H}_2\text{O})_2]$, the Zn^{II} cation, located on an inversion center, is N,N' -chelated by two 5-(pyrazin-2-yl)-3-(pyridin-3-yl)-1,2,4-triazolido anions and is further coordinated by two water molecules in a distorted N_4O_2 octahedral geometry. In the anionic ligand, the pyrazine and pyridine rings are twisted with respect to the central triazole ring by 5.77 (10) and 11.54 (10) $^\circ$, respectively. In the crystal, classical $\text{O}-\text{H}\cdots\text{N}$ and weak $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds and $\pi-\pi$ stacking interactions between aromatic rings [the centroid–centroid distances between triazole and pyrazine rings, and between triazole and pyridine rings are 3.623 (2) and 3.852 (2) \AA , respectively] connect the molecules into a three-dimensional supramolecular architecture.

Related literature

For applications and related structures of 1,2,4-triazole derivatives, see: Zhang *et al.* (2012); Chen *et al.* (2006).



Experimental

Crystal data

$[\text{Zn}(\text{C}_{11}\text{H}_7\text{N}_6)_2(\text{H}_2\text{O})_2]$

$M_r = 547.85$

Monoclinic, $P2_1/c$
 $a = 8.600$ (5) \AA
 $b = 5.728$ (3) \AA
 $c = 22.288$ (12) \AA
 $\beta = 100.646$ (6) $^\circ$
 $V = 1079.0$ (10) \AA^3

$Z = 2$
Mo $K\alpha$ radiation
 $\mu = 1.19\text{ mm}^{-1}$
 $T = 296\text{ K}$
 $0.18 \times 0.15 \times 0.13\text{ mm}$

Data collection

Bruker SMART 1000 CCD diffractometer
Absorption correction: multi-scan (*SADABS*; Bruker, 2001)
 $T_{\min} = 0.80$, $T_{\max} = 0.86$

11003 measured reflections
2493 independent reflections
2304 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.083$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.037$
 $wR(F^2) = 0.110$
 $S = 1.05$
2493 reflections
175 parameters
3 restraints

H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\max} = 0.44\text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.59\text{ e \AA}^{-3}$

Table 1
Selected bond lengths (\AA).

Zn1—O1	2.1054 (16)	Zn1—N5	2.1733 (16)
Zn1—N1	2.1853 (18)		

Table 2
Hydrogen-bond geometry (\AA , $^\circ$).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
O1—H1A \cdots N6 ⁱ	0.85 (2)	1.93 (2)	2.736 (2)	159 (2)
O1—H1B \cdots N4 ⁱⁱ	0.87 (2)	1.91 (2)	2.771 (2)	170 (2)
C1—H1 \cdots O1 ⁱⁱⁱ	0.93	2.41	3.266 (3)	153

Symmetry codes: (i) $x + 1, y + 1, z$; (ii) $-x, -y + 1, -z + 1$; (iii) $x, y + 1, z$.

Data collection: *SMART* (Bruker, 2007); cell refinement: *SAINT* (Bruker, 2007); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL*.

The authors are grateful to the Project of Hubei Provincial Education Office, China (grant No. Q20131304).

Supporting information for this paper is available from the IUCr electronic archives (Reference: XU5772).

References

- Bruker (2001). *SADABS*. Bruker AXS Inc., Madison, Wisconsin, USA.
- Bruker (2007). *SMART* and *SAINT*. Bruker AXS Inc., Madison, Wisconsin, USA.
- Chen, J.-C., Zhou, A.-J., Hu, S., Tong, M.-L. & Tong, Y.-X. (2006). *J. Mol. Struct.* **794**, 225–229.
- Sheldrick, G. M. (2008). *Acta Cryst. A* **64**, 112–122.
- Zhang, J.-P., Zhang, Y.-B., Lin, J.-B. & Chen, X.-M. (2012). *Chem. Rev.* **112**, 1001–1033.

supplementary materials

Acta Cryst. (2014). E70, m116 [doi:10.1107/S1600536814004176]

Diaquabis[5-(pyrazin-2-yl- κN^1)-3-(pyridin-3-yl)-1,2,4-triazolido- κN^1]zinc

Ye-Nan Wang and Wen-Wen Dong

1. Comment

1,2,4-Triazole derivatives are important building blocks of many important compounds widely used in medicine, agriculture, industry, and coordination chemistry (Zhang *et al.*, 2012; Chen *et al.*, 2006). During the synthesis of polymeric complexes using 2-(5-(pyridin-3-yl)-4H-1,2,4-triazol-3-yl)pyrazine as building blocks and, to our surprise, the title monomeric Zn(II) complex was obtained.

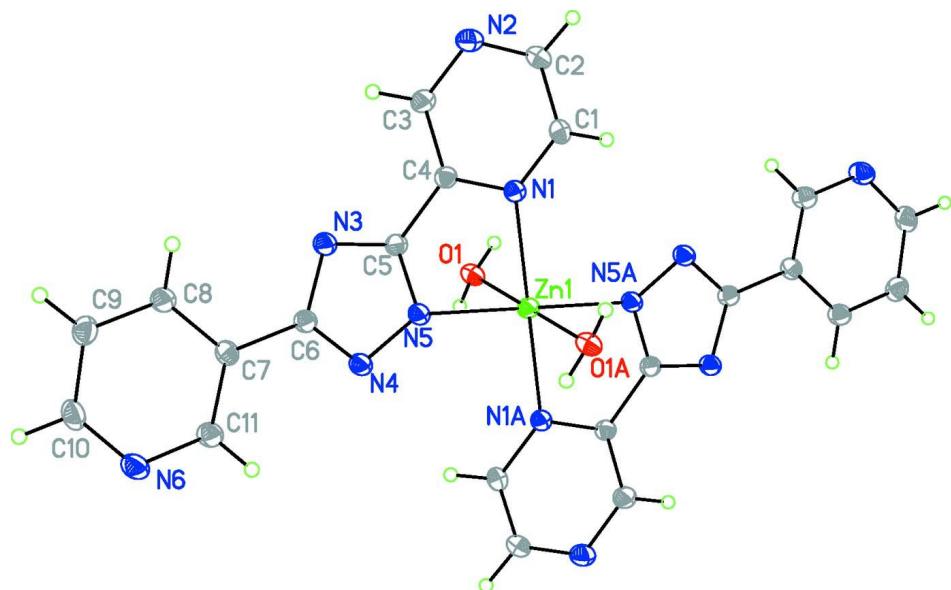
The title compound is a crystallographically centrosymmetric mononuclear complex. The Zn^{II} cation is in a distorted octahedral geometry (Fig. 1) and is coordinated by four N atoms from two 2-(5-(pyridin-3-yl)-1,2,4-triazolido-3-yl)pyrazine ligands and two coordinated water molecules. The observed Zn–O and Zn–N bond distances and bond angles reveal usual values. In the crystal, classic O–H \cdots N hydrogen bonds, weak C–H \cdots O hydrogen bond and π – π stacking between aromatic rings connect the molecules into the three dimensional supramolecular architecture [centroids between triazole and pyrazine rings and between triazole and pyridine rings being 3.623 (2) and 3.852 (2) Å, respectively].

2. Experimental

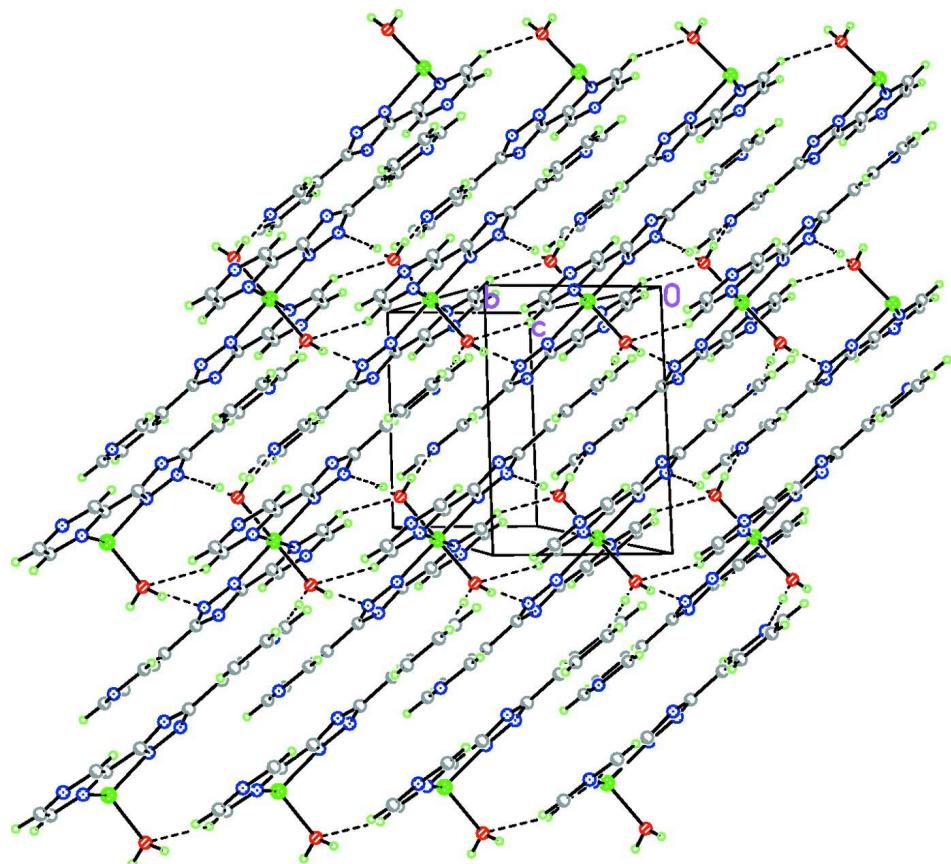
A mixture of 2-(5-(pyridin-3-yl)-4H-1,2,4-triazol-3-yl)pyrazine (0.0448 g, 0.2 mmol), Zn(CH₃COO)₂·2H₂O (0.0220 g, 0.1 mmol), water (6 mL), *N,N*-dimethylformamide (2 mL) was stirred vigorously for 30 min and then sealed in a Teflon-lined stainless-steel autoclave. The autoclave was heated and maintained at 433 K for 3 d, and then cooled to room temperature at 5 K h⁻¹ to obtain crystals suitable for X-ray analysis.

3. Refinement

All H-atoms were positioned geometrically and refined using a riding model with C–H = 0.93 Å, $U_{\text{iso}}(\text{H}) = 1.2 U_{\text{eq}}(\text{C})$ for aromatic hydrogen atoms. The H-atoms of O atoms were identified from a difference Fourier map and refined with O–H = 0.85 (2) Å, $U_{\text{iso}}(\text{H}) = 1.5 U_{\text{eq}}(\text{O})$.

**Figure 1**

The structure of the title compound showing the atomic numbering and 30% probability displacement ellipsoids.

**Figure 2**

A packing diagram.

Diaquabis[5-(pyrazin-2-yl- κN^1)-3-(pyridin-3-yl)-1,2,4-triazolido- κN^1]zinc*Crystal data* $[Zn(C_{11}H_7N_6)_2(H_2O)_2]$ $M_r = 547.85$ Monoclinic, $P2_1/c$

Hall symbol: -P 2ybc

 $a = 8.600 (5) \text{ \AA}$ $b = 5.728 (3) \text{ \AA}$ $c = 22.288 (12) \text{ \AA}$ $\beta = 100.646 (6)^\circ$ $V = 1079.0 (10) \text{ \AA}^3$ $Z = 2$ $F(000) = 560$ $D_x = 1.686 \text{ Mg m}^{-3}$ Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$

Cell parameters from 2880 reflections

 $\theta = 2.4-27.6^\circ$ $\mu = 1.19 \text{ mm}^{-1}$ $T = 296 \text{ K}$

Prism, colorless

 $0.18 \times 0.15 \times 0.13 \text{ mm}$ *Data collection*Bruker SMART 1000 CCD
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

phi and ω scansAbsorption correction: multi-scan
(SADABS; Bruker, 2001) $T_{\min} = 0.80, T_{\max} = 0.86$

11003 measured reflections

2493 independent reflections

2304 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.083$ $\theta_{\max} = 27.6^\circ, \theta_{\min} = 2.4^\circ$ $h = -11 \rightarrow 11$ $k = -7 \rightarrow 7$ $l = -29 \rightarrow 28$ *Refinement*Refinement on F^2

Least-squares matrix: full

 $R[F^2 > 2\sigma(F^2)] = 0.037$ $wR(F^2) = 0.110$ $S = 1.05$

2493 reflections

175 parameters

3 restraints

Primary atom site location: structure-invariant
direct methodsSecondary atom site location: difference Fourier
mapHydrogen site location: inferred from
neighbouring sitesH atoms treated by a mixture of independent
and constrained refinement $w = 1/[\sigma^2(F_o^2) + (0.0623P)^2 + 0.2938P]$
where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\max} < 0.001$ $\Delta\rho_{\max} = 0.44 \text{ e \AA}^{-3}$ $\Delta\rho_{\min} = -0.59 \text{ e \AA}^{-3}$ *Special details*

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R -factors(gt) etc. and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	x	y	z	$U_{\text{iso}}^* / U_{\text{eq}}$
Zn1	0.0000	1.0000	0.5000	0.02623 (14)
O1	0.17564 (15)	0.7938 (2)	0.47155 (6)	0.0298 (3)
H1B	0.218 (3)	0.679 (4)	0.4940 (11)	0.045*
H1A	0.252 (3)	0.871 (4)	0.4624 (11)	0.045*

N1	-0.03718 (18)	1.1583 (3)	0.40926 (7)	0.0255 (3)
N2	-0.0898 (2)	1.2803 (3)	0.28622 (8)	0.0364 (4)
N3	-0.32425 (18)	0.7016 (3)	0.35570 (7)	0.0273 (3)
N4	-0.27382 (18)	0.5864 (3)	0.45419 (7)	0.0271 (3)
N5	-0.17912 (18)	0.7731 (3)	0.44860 (7)	0.0258 (3)
N6	-0.6262 (2)	0.0459 (3)	0.41338 (9)	0.0357 (4)
C1	0.0312 (2)	1.3446 (3)	0.38951 (9)	0.0302 (4)
H1	0.0992	1.4355	0.4174	0.036*
C2	0.0033 (3)	1.4058 (3)	0.32853 (9)	0.0339 (4)
H2	0.0512	1.5393	0.3167	0.041*
C3	-0.1585 (3)	1.0954 (4)	0.30615 (9)	0.0331 (4)
H3	-0.2251	1.0039	0.2779	0.040*
C4	-0.1352 (2)	1.0330 (3)	0.36728 (9)	0.0238 (4)
C5	-0.2139 (2)	0.8348 (3)	0.38977 (8)	0.0242 (4)
C6	-0.3581 (2)	0.5502 (3)	0.39810 (9)	0.0249 (4)
C7	-0.4734 (2)	0.3611 (3)	0.38268 (8)	0.0261 (4)
C8	-0.5332 (2)	0.3074 (4)	0.32153 (9)	0.0312 (4)
H8	-0.5029	0.3951	0.2905	0.037*
C9	-0.6372 (2)	0.1240 (4)	0.30748 (10)	0.0353 (4)
H9	-0.6783	0.0865	0.2671	0.042*
C10	-0.6790 (3)	-0.0028 (3)	0.35465 (12)	0.0362 (5)
H10	-0.7475	-0.1285	0.3450	0.043*
C11	-0.5265 (2)	0.2270 (4)	0.42666 (9)	0.0317 (4)
H11	-0.4912	0.2645	0.4675	0.038*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Zn1	0.0291 (2)	0.0287 (2)	0.0197 (2)	-0.00523 (11)	0.00119 (13)	0.00147 (10)
O1	0.0282 (7)	0.0286 (7)	0.0331 (7)	-0.0035 (5)	0.0064 (5)	0.0056 (5)
N1	0.0268 (7)	0.0255 (7)	0.0238 (7)	-0.0033 (6)	0.0040 (6)	0.0002 (5)
N2	0.0442 (10)	0.0369 (9)	0.0274 (8)	-0.0051 (8)	0.0044 (7)	0.0068 (7)
N3	0.0269 (7)	0.0293 (8)	0.0249 (7)	-0.0060 (6)	0.0023 (6)	0.0017 (6)
N4	0.0266 (7)	0.0279 (8)	0.0264 (8)	-0.0056 (6)	0.0042 (6)	0.0020 (6)
N5	0.0273 (7)	0.0253 (7)	0.0240 (7)	-0.0057 (6)	0.0030 (6)	0.0013 (6)
N6	0.0303 (9)	0.0395 (9)	0.0382 (10)	-0.0096 (7)	0.0089 (8)	0.0040 (8)
C1	0.0326 (9)	0.0266 (9)	0.0314 (9)	-0.0077 (7)	0.0059 (7)	-0.0027 (7)
C2	0.0414 (11)	0.0271 (9)	0.0344 (10)	-0.0045 (8)	0.0105 (8)	0.0050 (8)
C3	0.0386 (10)	0.0359 (10)	0.0234 (9)	-0.0081 (8)	0.0018 (7)	0.0009 (8)
C4	0.0229 (8)	0.0257 (8)	0.0227 (9)	-0.0007 (6)	0.0036 (7)	0.0005 (6)
C5	0.0237 (8)	0.0266 (9)	0.0222 (8)	-0.0032 (7)	0.0036 (6)	0.0002 (6)
C6	0.0212 (8)	0.0289 (8)	0.0246 (9)	-0.0035 (7)	0.0041 (7)	0.0005 (7)
C7	0.0203 (8)	0.0277 (8)	0.0300 (9)	-0.0035 (7)	0.0040 (7)	-0.0002 (7)
C8	0.0295 (9)	0.0353 (10)	0.0281 (9)	-0.0078 (8)	0.0040 (7)	0.0013 (8)
C9	0.0318 (10)	0.0409 (11)	0.0324 (10)	-0.0073 (8)	0.0036 (8)	-0.0064 (8)
C10	0.0306 (11)	0.0336 (12)	0.0442 (14)	-0.0114 (7)	0.0063 (10)	-0.0056 (8)
C11	0.0250 (8)	0.0396 (10)	0.0304 (9)	-0.0077 (8)	0.0048 (7)	0.0014 (8)

Geometric parameters (\AA , ^\circ)

Zn1—O1 ⁱ	2.1054 (16)	N6—C10	1.334 (3)
Zn1—O1	2.1054 (16)	N6—C11	1.344 (3)
Zn1—N1	2.1853 (18)	C1—C2	1.381 (3)
Zn1—N1 ⁱ	2.1853 (18)	C1—H1	0.9300
Zn1—N5	2.1733 (16)	C2—H2	0.9300
Zn1—N5 ⁱ	2.1733 (16)	C3—C4	1.387 (3)
O1—H1B	0.867 (16)	C3—H3	0.9300
O1—H1A	0.848 (16)	C4—C5	1.457 (2)
N1—C1	1.332 (2)	C6—C7	1.466 (3)
N1—C4	1.345 (2)	C7—C11	1.387 (3)
N2—C3	1.329 (3)	C7—C8	1.399 (3)
N2—C2	1.330 (3)	C8—C9	1.377 (3)
N3—C5	1.339 (2)	C8—H8	0.9300
N3—C6	1.353 (2)	C9—C10	1.379 (3)
N4—C6	1.340 (2)	C9—H9	0.9300
N4—N5	1.364 (2)	C10—H10	0.9300
N5—C5	1.338 (2)	C11—H11	0.9300
O1 ⁱ —Zn1—O1	180.00 (7)	N2—C2—C1	122.23 (18)
O1 ⁱ —Zn1—N5	90.93 (7)	N2—C2—H2	118.9
O1—Zn1—N5	89.07 (7)	C1—C2—H2	118.9
O1 ⁱ —Zn1—N5 ⁱ	89.07 (7)	N2—C3—C4	122.76 (18)
O1—Zn1—N5 ⁱ	90.93 (7)	N2—C3—H3	118.6
N5—Zn1—N5 ⁱ	180.0	C4—C3—H3	118.6
O1 ⁱ —Zn1—N1	93.18 (6)	N1—C4—C3	120.28 (17)
O1—Zn1—N1	86.82 (6)	N1—C4—C5	116.58 (17)
N5—Zn1—N1	78.00 (6)	C3—C4—C5	123.13 (17)
N5 ⁱ —Zn1—N1	102.00 (6)	N5—C5—N3	114.31 (16)
O1 ⁱ —Zn1—N1 ⁱ	86.82 (6)	N5—C5—C4	120.63 (16)
O1—Zn1—N1 ⁱ	93.18 (6)	N3—C5—C4	125.06 (16)
N5—Zn1—N1 ⁱ	102.00 (6)	N4—C6—N3	113.79 (17)
N5 ⁱ —Zn1—N1 ⁱ	78.00 (6)	N4—C6—C7	123.96 (17)
N1—Zn1—N1 ⁱ	180.000 (1)	N3—C6—C7	122.22 (17)
Zn1—O1—H1B	120.2 (18)	C11—C7—C8	117.17 (17)
Zn1—O1—H1A	114.1 (18)	C11—C7—C6	122.69 (17)
H1B—O1—H1A	106 (2)	C8—C7—C6	120.14 (17)
C1—N1—C4	117.11 (16)	C9—C8—C7	119.71 (18)
C1—N1—Zn1	129.81 (13)	C9—C8—H8	120.1
C4—N1—Zn1	112.83 (12)	C7—C8—H8	120.1
C3—N2—C2	116.14 (17)	C8—C9—C10	118.57 (19)
C5—N3—C6	100.99 (15)	C8—C9—H9	120.7
C6—N4—N5	105.37 (15)	C10—C9—H9	120.7
C5—N5—N4	105.55 (14)	N6—C10—C9	123.25 (19)
C5—N5—Zn1	111.52 (11)	N6—C10—H10	118.4
N4—N5—Zn1	142.93 (12)	C9—C10—H10	118.4
C10—N6—C11	117.76 (19)	N6—C11—C7	123.48 (19)
N1—C1—C2	121.43 (18)	N6—C11—H11	118.3

N1—C1—H1	119.3	C7—C11—H11	118.3
C2—C1—H1	119.3		

Symmetry code: (i) $-x, -y+2, -z+1$.

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

$D\text{—H}\cdots A$	$D\text{—H}$	$H\cdots A$	$D\cdots A$	$D\text{—H}\cdots A$
O1—H1A \cdots N6 ⁱⁱ	0.85 (2)	1.93 (2)	2.736 (2)	159 (2)
O1—H1B \cdots N4 ⁱⁱⁱ	0.87 (2)	1.91 (2)	2.771 (2)	170 (2)
C1—H1 \cdots O1 ^{iv}	0.93	2.41	3.266 (3)	153

Symmetry codes: (ii) $x+1, y+1, z$; (iii) $-x, -y+1, -z+1$; (iv) $x, y+1, z$.