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Tetraaquabis[2-[4-(4-pyridyl)pyrimidin-2-ylsulfanyl]acetato]zinc

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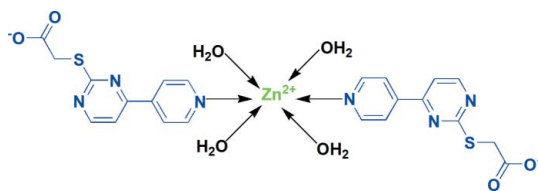
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 Key indicators: single-crystal X-ray study; $T = 298$ K; mean $\sigma(\text{C}-\text{C}) = 0.004$ Å; R factor = 0.033; wR factor = 0.086; data-to-parameter ratio = 13.0.

In the title compound, $[\text{Zn}(\text{C}_{11}\text{H}_8\text{N}_3\text{O}_2\text{S})_2(\text{H}_2\text{O})_4]$, the Zn^{II} ion lies on an inversion centre and is coordinated by four water molecules and two N atoms from two 2-[4-(4-pyridyl)pyrimidin-2-ylsulfanyl]acetate (*L*) ligands in a distorted octahedral geometry. In *L*, the pyridine and pyrimidine rings are twisted at an angle of $11.2(1)^\circ$. The coordinated water molecules and the acetate groups are involved in the formation of a three-dimensional hydrogen-bonded network, which consolidates the crystal packing.

Related literature

 For a related structure, see: Zhu *et al.* (2009).


Experimental

Crystal data

 $[\text{Zn}(\text{C}_{11}\text{H}_8\text{N}_3\text{O}_2\text{S})_2(\text{H}_2\text{O})_4]$
 $M_r = 630.00$
 Orthorhombic, *Pbca*
 $a = 7.199(7)$ Å
 $b = 11.792(11)$ Å
 $c = 28.77(3)$ Å

 $V = 2442(4)$ Å³
 $Z = 4$
 Mo $K\alpha$ radiation

 $\mu = 1.24$ mm⁻¹
 $T = 298$ K
 $0.20 \times 0.20 \times 0.15$ mm

Data collection

 Bruker APEXII CCD area-detector diffractometer
 Absorption correction: multi-scan (*SADABS*; Bruker, 2001)
 $T_{\text{min}} = 0.780$, $T_{\text{max}} = 0.830$

 16776 measured reflections
 2465 independent reflections
 1836 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.051$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.033$
 $wR(F^2) = 0.086$
 $S = 1.02$
 2465 reflections
 190 parameters
 4 restraints

 H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\text{max}} = 0.23$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.45$ e Å⁻³
Table 1

Hydrogen-bond geometry (Å, °).

<i>D</i> —H··· <i>A</i>	<i>D</i> —H	H··· <i>A</i>	<i>D</i> ··· <i>A</i>	<i>D</i> —H··· <i>A</i>
O4—H4WB···O2 ⁱ	0.81 (2)	1.96 (2)	2.759 (3)	170 (3)
O4—H4WA···O1 ⁱⁱ	0.82 (2)	1.82 (2)	2.632 (3)	172 (2)
O3—H3WB···O2 ⁱⁱⁱ	0.84 (2)	1.89 (2)	2.728 (3)	176 (3)
O3—H3WA···O2 ^{iv}	0.82 (2)	2.24 (2)	3.060 (3)	172 (3)

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

Data collection: *APEX2* (Bruker, 2007); cell refinement: *SAINT-Plus* (Bruker, 2007); data reduction: *SAINT-Plus*; 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*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: CV5136).

References

- Bruker (2001). *SADABS*. Bruker AXS Inc., Madison, Wisconsin, USA.
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 Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112–122.
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supporting information

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Tetraaquabis{2-[4-(4-pyridyl)pyrimidin-2-ylsulfanyl]acetato}zinc

Hai-Bin Zhu and Xin Lu

S1. Comment

In our previous work, we have reported the crystal structure of mononuclear Mn(II) complex with the ligand of 2-(4-(pyridine-3-yl)pyrimidin-2-ylthio)acetic acid (Zhu *et al.*, 2009). Herein, we present a Zn(II) complex with the ligand of 2-(4-(pyridine-4-yl)pyrimidin-2-ylthio)acetic acid.

Similar to the reported Mn(II) coordination compound (Zhu *et al.*, 2009), the Zn(II) center in the title compound also adopts an octahedral coordination geometry defined by four water O atoms in equatorial positions and two N atoms in apical positions (Fig. 1). The Zn—O bond lengths vary from 2.070 (2) to 2.137 (2) Å, and the Zn—N bond length is 2.176 (2) Å. Intermolecular O—H...O hydrogen bonds (Table 1) consolidate the crystal packing.

S2. Experimental

The mixture of Zn(NO₃)₂ (0.1 mmol), *L* (0.2 mmol) and NaOH (0.2 mmol) in 10 ml of H₂O was stirred for 30 min at room temperature. After filtration, the mother liquid was stood for three weeks to give yellow crystals suitable for X-ray diffraction analysis.

S3. Refinement

C-bound H atoms were positioned geometrically (C—H = 0.93 Å) and allowed to ride on their parent atoms, with $U_{\text{iso}}(\text{H}) = 1.2 U_{\text{eq}}(\text{C})$. The positions of the water H atoms were found from a difference Fourier map and refined with restraint O—H = 0.82 (2) Å using a riding model, with $U_{\text{iso}}(\text{H}) = 1.2 U_{\text{eq}}(\text{O})$.

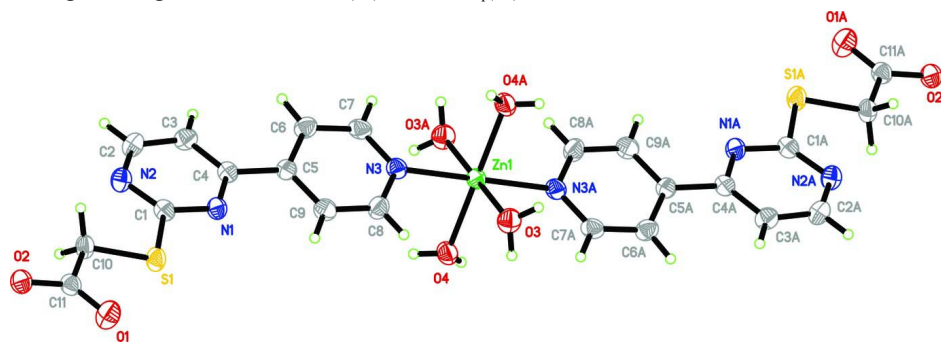


Figure 1

The coordination environment around Zn(II) in the title complex with the atom-labeling scheme [symmetry code: (A) $-x, -y + 1, -z + 1$]. Displacement ellipsoids for non-hydrogen atoms are drawn at the 50% probability level.

Tetraaquabis{2-[4-(4-pyridyl)pyrimidin-2-ylsulfanyl]acetato}zinc

Crystal data

[Zn(C₁₁H₈N₃O₂S)₂(H₂O)₄] $M_r = 630.00$ Orthorhombic, *Pbca*

Hall symbol: -P 2ac 2ab

 $a = 7.199$ (7) Å $b = 11.792$ (11) Å $c = 28.77$ (3) Å $V = 2442$ (4) Å³ $Z = 4$ $F(000) = 1296$ $D_x = 1.714$ Mg m⁻³Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 2465 reflections

 $\theta = 2.3$ – 25.5° $\mu = 1.24$ mm⁻¹ $T = 298$ K

Block, yellow

 $0.20 \times 0.20 \times 0.15$ mm

Data collection

Bruker APEXII CCD area-detector
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

 φ and ω scansAbsorption correction: multi-scan
(*SADABS*; Bruker, 2001) $T_{\min} = 0.780$, $T_{\max} = 0.830$

16776 measured reflections

2465 independent reflections

1836 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.051$ $\theta_{\max} = 27.1^\circ$, $\theta_{\min} = 1.4^\circ$ $h = -8 \rightarrow 9$ $k = -14 \rightarrow 14$ $l = -34 \rightarrow 32$

Refinement

Refinement on F^2

Least-squares matrix: full

 $R[F^2 > 2\sigma(F^2)] = 0.033$ $wR(F^2) = 0.086$ $S = 1.02$

2465 reflections

190 parameters

4 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 $w = 1/[\sigma^2(F_o^2) + (0.0432P)^2 + 0.3573P]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\max} < 0.001$ $\Delta\rho_{\max} = 0.23$ e Å⁻³ $\Delta\rho_{\min} = -0.45$ e Å⁻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 (Å²)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
Zn1	0.0000	0.5000	0.5000	0.02474 (14)
S1	0.01224 (8)	0.60147 (5)	0.17445 (2)	0.02955 (17)
N2	0.1353 (3)	0.80094 (16)	0.20795 (6)	0.0297 (5)
N1	0.0402 (3)	0.65524 (16)	0.26077 (6)	0.0264 (5)

O4	0.2360 (2)	0.41441 (14)	0.47861 (5)	0.0314 (4)
H4WB	0.307 (3)	0.396 (2)	0.4993 (7)	0.038*
H4WA	0.302 (3)	0.450 (2)	0.4603 (7)	0.038*
C1	0.0700 (3)	0.6981 (2)	0.21845 (8)	0.0256 (5)
C10	0.0382 (3)	0.6859 (2)	0.12255 (8)	0.0283 (5)
H10A	-0.0412	0.7522	0.1244	0.034*
H10B	0.1658	0.7114	0.1198	0.034*
O1	-0.0740 (3)	0.51902 (14)	0.08678 (6)	0.0396 (5)
C5	0.0435 (3)	0.67551 (18)	0.34366 (8)	0.0236 (5)
C4	0.0801 (3)	0.72292 (19)	0.29679 (7)	0.0241 (5)
C11	-0.0136 (3)	0.6163 (2)	0.08050 (8)	0.0269 (5)
N3	-0.0130 (2)	0.58138 (16)	0.43217 (6)	0.0255 (4)
O2	0.0038 (2)	0.66402 (14)	0.04128 (6)	0.0339 (4)
O3	-0.1746 (3)	0.36656 (14)	0.47517 (6)	0.0342 (4)
H3WB	-0.118 (3)	0.3056 (17)	0.4696 (9)	0.041*
H3WA	-0.269 (3)	0.355 (2)	0.4908 (8)	0.041*
C7	0.0205 (3)	0.6920 (2)	0.42625 (8)	0.0287 (6)
H7A	0.0259	0.7377	0.4525	0.034*
C3	0.1516 (3)	0.83095 (19)	0.28983 (8)	0.0295 (5)
H3B	0.1814	0.8781	0.3147	0.035*
C9	0.0026 (3)	0.5610 (2)	0.34959 (8)	0.0301 (6)
H9A	-0.0059	0.5135	0.3239	0.036*
C8	-0.0249 (3)	0.5184 (2)	0.39326 (8)	0.0309 (6)
H8A	-0.0534	0.4418	0.3962	0.037*
C2	0.1763 (4)	0.86515 (19)	0.24447 (8)	0.0311 (6)
H2B	0.2245	0.9372	0.2391	0.037*
C6	0.0476 (3)	0.7421 (2)	0.38362 (8)	0.0294 (5)
H6A	0.0685	0.8197	0.3815	0.035*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Zn1	0.0297 (2)	0.0257 (2)	0.0188 (2)	0.00061 (16)	0.00071 (16)	-0.00053 (15)
S1	0.0392 (4)	0.0303 (3)	0.0191 (3)	-0.0049 (3)	-0.0001 (3)	0.0019 (2)
N2	0.0318 (11)	0.0314 (11)	0.0259 (10)	-0.0026 (9)	-0.0017 (9)	0.0039 (9)
N1	0.0317 (12)	0.0287 (11)	0.0187 (10)	0.0006 (8)	-0.0007 (8)	0.0016 (8)
O4	0.0326 (10)	0.0363 (10)	0.0255 (9)	0.0049 (8)	0.0041 (8)	0.0036 (8)
C1	0.0233 (12)	0.0312 (13)	0.0222 (12)	0.0020 (10)	-0.0017 (9)	-0.0006 (10)
C10	0.0328 (14)	0.0304 (13)	0.0216 (12)	-0.0034 (10)	0.0017 (10)	0.0032 (10)
O1	0.0562 (12)	0.0353 (10)	0.0274 (10)	-0.0098 (9)	-0.0089 (9)	0.0006 (8)
C5	0.0238 (12)	0.0261 (12)	0.0210 (12)	0.0020 (9)	-0.0029 (9)	0.0013 (10)
C4	0.0218 (12)	0.0287 (12)	0.0217 (12)	0.0019 (10)	-0.0018 (10)	0.0008 (10)
C11	0.0275 (13)	0.0307 (13)	0.0226 (12)	0.0033 (10)	0.0000 (10)	0.0015 (10)
N3	0.0291 (11)	0.0275 (11)	0.0200 (10)	0.0010 (8)	-0.0004 (8)	-0.0019 (8)
O2	0.0474 (11)	0.0340 (10)	0.0203 (9)	0.0031 (8)	0.0014 (7)	0.0017 (7)
O3	0.0373 (11)	0.0329 (10)	0.0325 (10)	-0.0045 (8)	-0.0010 (8)	-0.0016 (8)
C7	0.0347 (14)	0.0288 (13)	0.0225 (12)	0.0031 (10)	-0.0018 (10)	-0.0062 (10)
C3	0.0339 (14)	0.0271 (13)	0.0276 (13)	-0.0002 (10)	-0.0047 (11)	-0.0022 (10)

C9	0.0393 (15)	0.0289 (13)	0.0220 (12)	-0.0023 (11)	-0.0014 (10)	-0.0066 (10)
C8	0.0436 (16)	0.0257 (12)	0.0233 (13)	-0.0046 (11)	-0.0005 (11)	-0.0001 (10)
C2	0.0328 (14)	0.0267 (13)	0.0339 (14)	-0.0042 (11)	-0.0028 (11)	0.0045 (11)
C6	0.0358 (14)	0.0247 (12)	0.0277 (13)	0.0001 (10)	-0.0036 (11)	-0.0015 (10)

Geometric parameters (Å, °)

Zn1—O4	2.070 (2)	C5—C9	1.393 (3)
Zn1—O4 ⁱ	2.070 (2)	C5—C6	1.392 (3)
Zn1—O3 ⁱ	2.137 (2)	C5—C4	1.483 (3)
Zn1—O3	2.137 (2)	C4—C3	1.388 (3)
Zn1—N3 ⁱ	2.176 (2)	C11—O2	1.267 (3)
Zn1—N3	2.176 (2)	N3—C7	1.337 (3)
S1—C1	1.753 (3)	N3—C8	1.346 (3)
S1—C10	1.804 (3)	O3—H3WB	0.843 (16)
N2—C2	1.328 (3)	O3—H3WA	0.822 (17)
N2—C1	1.335 (3)	C7—C6	1.375 (3)
N1—C1	1.336 (3)	C7—H7A	0.9300
N1—C4	1.339 (3)	C3—C2	1.377 (3)
O4—H4WB	0.810 (16)	C3—H3B	0.9300
O4—H4WA	0.821 (16)	C9—C8	1.367 (4)
C10—C11	1.508 (3)	C9—H9A	0.9300
C10—H10A	0.9700	C8—H8A	0.9300
C10—H10B	0.9700	C2—H2B	0.9300
O1—C11	1.240 (3)	C6—H6A	0.9300
O4—Zn1—O4 ⁱ	180.00 (8)	C6—C5—C4	122.3 (2)
O4—Zn1—O3 ⁱ	88.60 (9)	N1—C4—C3	121.0 (2)
O4 ⁱ —Zn1—O3 ⁱ	91.40 (9)	N1—C4—C5	116.1 (2)
O4—Zn1—O3	91.40 (9)	C3—C4—C5	122.9 (2)
O4 ⁱ —Zn1—O3	88.60 (9)	O1—C11—O2	125.1 (2)
O3 ⁱ —Zn1—O3	180.0	O1—C11—C10	118.2 (2)
O4—Zn1—N3 ⁱ	90.94 (7)	O2—C11—C10	116.6 (2)
O4 ⁱ —Zn1—N3 ⁱ	89.06 (7)	C7—N3—C8	116.3 (2)
O3 ⁱ —Zn1—N3 ⁱ	89.99 (8)	C7—N3—Zn1	122.44 (15)
O3—Zn1—N3 ⁱ	90.01 (8)	C8—N3—Zn1	120.33 (17)
O4—Zn1—N3	89.06 (7)	Zn1—O3—H3WB	113.9 (19)
O4 ⁱ —Zn1—N3	90.94 (7)	Zn1—O3—H3WA	114.9 (19)
O3 ⁱ —Zn1—N3	90.01 (8)	H3WB—O3—H3WA	111 (3)
O3—Zn1—N3	89.99 (8)	N3—C7—C6	123.9 (2)
N3 ⁱ —Zn1—N3	180.0	N3—C7—H7A	118.0
C1—S1—C10	102.37 (13)	C6—C7—H7A	118.0
C2—N2—C1	114.66 (19)	C2—C3—C4	116.9 (2)
C1—N1—C4	116.4 (2)	C2—C3—H3B	121.5
Zn1—O4—H4WB	115.2 (19)	C4—C3—H3B	121.5
Zn1—O4—H4WA	114.8 (18)	C8—C9—C5	119.9 (2)
H4WB—O4—H4WA	104 (3)	C8—C9—H9A	120.0
N2—C1—N1	127.3 (2)	C5—C9—H9A	120.0

N2—C1—S1	120.71 (17)	N3—C8—C9	123.5 (2)
N1—C1—S1	111.96 (18)	N3—C8—H8A	118.2
C11—C10—S1	109.75 (17)	C9—C8—H8A	118.2
C11—C10—H10A	109.7	N2—C2—C3	123.6 (2)
S1—C10—H10A	109.7	N2—C2—H2B	118.2
C11—C10—H10B	109.7	C3—C2—H2B	118.2
S1—C10—H10B	109.7	C7—C6—C5	119.4 (2)
H10A—C10—H10B	108.2	C7—C6—H6A	120.3
C9—C5—C6	116.8 (2)	C5—C6—H6A	120.3
C9—C5—C4	121.0 (2)		

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

Hydrogen-bond geometry (Å, °)

<i>D—H...A</i>	<i>D—H</i>	<i>H...A</i>	<i>D...A</i>	<i>D—H...A</i>
O4—H4WB...O2 ⁱⁱ	0.81 (2)	1.96 (2)	2.759 (3)	170 (3)
O4—H4WA...O1 ⁱⁱⁱ	0.82 (2)	1.82 (2)	2.632 (3)	172 (2)
O3—H3WB...O2 ^{iv}	0.84 (2)	1.89 (2)	2.728 (3)	176 (3)
O3—H3WA...O2 ^v	0.82 (2)	2.24 (2)	3.060 (3)	172 (3)

Symmetry codes: (ii) $-x+1/2, -y+1, z+1/2$; (iii) $x+1/2, y, -z+1/2$; (iv) $-x, y-1/2, -z+1/2$; (v) $-x-1/2, -y+1, z+1/2$.