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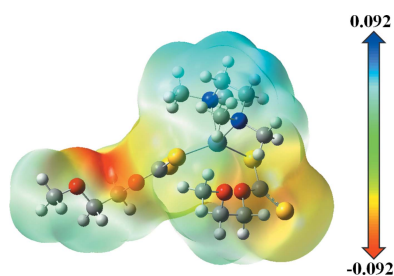
Crystal structure and DFT study of a zinc xanthate complex

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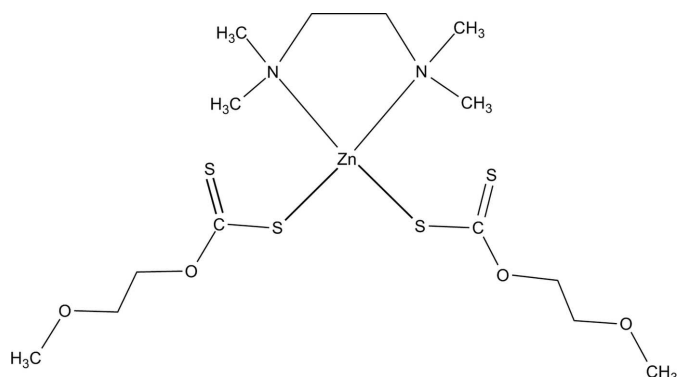
In the title compound, bis(2-methoxyethyl xanthato- κ S)(*N,N,N',N'*-tetramethylethylenediamine- κ^2N,N')zinc(II) acetone hemisolvate, $[\text{Zn}(\text{C}_4\text{H}_7\text{O}_2\text{S}_2)_2(\text{C}_6\text{H}_{16}\text{N}_2)] \cdot 0.5\text{C}_3\text{H}_6\text{O}$, the Zn^{II} ion is coordinated by two N atoms of the *N,N,N',N'*-tetramethylethylenediamine ligand and two S atoms from two 2-methoxyethyl xanthate ligands. The amine ligand is disordered over two orientations and was modelled with refined occupancies of 0.538 (6) and 0.462 (6). The molecular structure features two C—H \cdots O and two C—H \cdots S intramolecular interactions. In the crystal, molecules are linked by weak C—H \cdots O and C—H \cdots S hydrogen bonds, forming a three-dimensional supramolecular architecture. The molecular structure was optimized using density functional theory (DFT) at the B3LYP/6-311 G(d,p) level. The smallest HOMO–LUMO energy gap (3.19 eV) indicates the suitability of this crystal for optoelectronic applications. The molecular electrostatic potential (MEP) further identifies the positive, negative and neutral electrostatic potential regions of the molecules. Half a molecule of disordered acetone was removed with the solvent-mask procedure in *OLEX2* [Dolomanov *et al.* (2009)]. *J. Appl. Cryst.* **42**, 339–341] and this contribution is included in the formula.

1. Chemical context

Xanthates (dithiocarbonates) are related to the dithiolate family. Xanthate is a bidentate monoanionic sulfur–sulfur donor ligand. It stabilizes complexes of most of the transition elements and can bind metal centers in monodentate, isobidentate, anisobidentate or ionic modes. Xanthates have the ability to inhibit the replication of both RNA and DNA viruses *in vitro* (Friebolin *et al.*, 2005). They have been used as accelerators in the vulcanization of rubber (Gupta *et al.*, 2012), in cellulose synthesis (Tiravanti *et al.*, 1998), as collectors in the froth flotation of metal sulfide ores (Lee *et al.*, 2009) and as reagents for heavy-metal sedimentation in waste-water treatment (Chakraborty *et al.*, 2006). In our previous work, we prepared and structurally characterized nickel(II) and zinc(II) *n*-propylxanthate complexes containing *N,N,N',N'*-tetramethylethylenediamine as a neutral ligand. Both complexes showed a distorted octahedral environment around the metal center (Qadir & Dege, 2019). In this paper, we report the synthesis and crystal structure of a zinc(II) 2-methoxyethylxanthate complex containing *N,N,N',N'*-tetramethylethylenediamine, $[\text{Zn}(\text{S}_2\text{COC}_2\text{H}_4\text{OCH}_3)_2(\text{tmeda})]$, which was investigated by a DFT study.



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2. Structural commentary

The molecular structure of the title compound is illustrated in Fig. 1. The Zn^{II} ion is coordinated by two N atoms of the *N,N,N,N*-tetramethylethylenediamine molecule and two S atoms from two 2-methoxyethylxanthate molecules. The Zn1–N1, Zn1–N2, Zn1–S1 and Zn1–S3 bond lengths are 2.141 (5), 2.123 (5), 2.3107 (9) and 2.3050 (9) Å, respectively (Table 1). These bond distances are similar to those reported in the work of Cusack *et al.* (2004). The C7–O8 and C13–O14 bond lengths are similar [1.344 (3) and 1.346 (3) Å, respectively], while the C9–O8 and C15–O14 bonds are also not significantly different [1.454 (3) and 1.459 (3) Å, respectively]. In the same way, the C10–O11 [1.417 (3)] and C16–O17 [1.418 (4)] bond lengths are similar to each other. All of

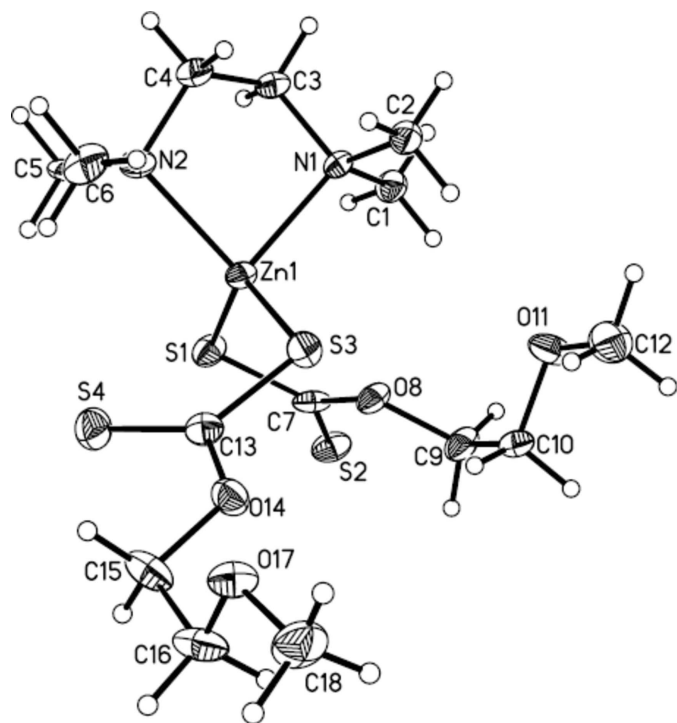


Figure 1
The molecular structure of the title complex, with the atom labelling. Only the major component of the disordered amine ligand is shown. Displacement ellipsoids are drawn at the 50% probability level.

Table 1
Selected geometric parameters (Å, °).

Zn1–S1	2.3107 (9)	S1–C7	1.731 (3)
Zn1–S3	2.3050 (9)	S2–C7	1.647 (3)
Zn1–N1	2.141 (5)	S3–C13	1.723 (3)
Zn1–N2	2.123 (5)	S4–C13	1.657 (3)
S3–Zn1–S1	125.54 (3)	N1–Zn1–S3	106.5 (2)
N1–Zn1–S1	105.2 (2)	N2–Zn1–N1	86.9 (2)

Table 2
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>
C1–H1C···O8	0.98	2.48	3.103 (7)	121
C2A–H2AB···O11	0.98	2.24	3.207 (13)	168
C5A–H5AA···S1	0.98	2.92	3.454 (16)	115
C6–H6C···S4	0.98	2.74	3.512 (13)	136
C6–H6B···O11 ⁱ	0.98	2.54	3.321 (13)	136
C3A–H3AB···S2 ⁱⁱ	0.99	2.81	3.483 (7)	125
C6A–H6AA···S1 ⁱⁱ	0.98	2.84	3.764 (16)	158
C4A–H4AA···O17 ⁱⁱⁱ	0.99	2.44	3.380 (6)	159
C4A–H4AB···S3 ⁱⁱⁱ	0.99	2.81	3.774 (8)	164
C9–H9A···O17 ^{iv}	0.99	2.61	3.415 (4)	138
C9–H9B···S2 ^v	0.99	2.94	3.708 (3)	135
C18–H18B···S2 ^{vi}	0.98	3.02	3.998 (3)	176

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

the C–O bonds show single-bond character. In the {S₂C} section of the xanthate ligands, the carbon-to-sulfur S1 distance is 1.731 (3) Å, which is typical of a single bond whereas the carbon-to-sulfur S2 distance of 1.647 (3) Å is typical of a carbon-to-sulfur double bond. In the molecule, weak C1–H1C···O8, C2A–H2AB···O11, C5A–H5AA···S1 and C6–H6C···S4 intramolecular interactions are observed (Table 2).

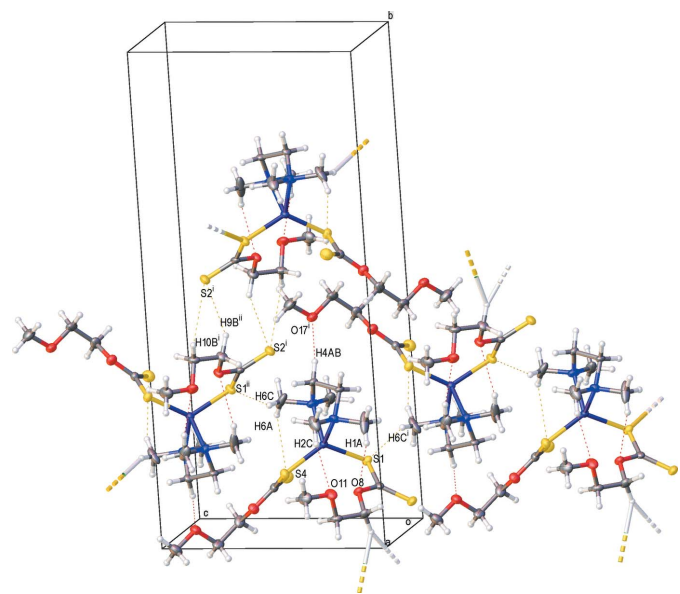


Figure 2
A view of the crystal packing of the title complex. Dashed lines denote the intermolecular hydrogen bonds (Table 2). Symmetry codes: (i) $x - \frac{1}{2}, -y + \frac{1}{2}, z - \frac{1}{2}$; (ii) $x + \frac{1}{2}, -y + \frac{1}{2}, z + \frac{1}{2}$.

3. Supramolecular features

The crystal packing of the title compound (Fig. 2) features intermolecular hydrogen bonds (C6–H6B···O11ⁱ, C3A–H3AB···S2ⁱⁱ, C6A–H6AA···S1ⁱⁱ, C4A–H4AA···O17ⁱⁱⁱ, C4A–H4AB···S3ⁱⁱⁱ, C9–H9A···O17^{iv}, C9–H9B···S2^v and C18–H18B···S2^{vi}; symmetry codes as in Table 2), which connect the molecules into a three-dimensional supramolecular architecture.

4. Database survey

Previously reported complexes related to the title complex are [Cd(S₂COCH₂CH₂OMe)₂(bipy)] [CSD (Groom *et al.*, 2016) refcode BENDII; Chen *et al.*, 2002], [Ni(C₄H₇O₂S₂)₂(C₆H₁₆N₂)] (NADTAQ; Qadir, 2016), [Ni(moexa)₂phen] (unsolvated form) and [Ni(moexa)₂phen] (benzene solvate), moexa = *O*-methoxyethylxanthato-*S,S'* (with refcodes SICTUT and SICVAB, respectively; Edwards *et al.*, 1990a), [Ni(moexa)₂bpy]; forms I and II (with refcodes VETVIZ and VETVIZ01, respectively; Edwards *et al.*, 1990b) and [Cd(S₂COCH₂CH₂OCH₃)₂(4,7-Me₂phen)] (WACPOG; Chen *et al.*, 2003). The Cd–S and Cd–N bond lengths range from 2.489 to 2.796 Å and 2.334 to 2.406 Å, respectively. Similarly, the Ni–S and Ni–N bond lengths range from 2.432 to 2.458 Å and 2.070 to 2.172 Å, respectively. In these complexes, compared with the Zn^{II} complex, the metal-to-ligand distances with *M*–S/N bond lengths follow the order Zn^{II} < Ni^{II} < Cd^{II} in the corresponding complexes.

5. Frontier molecular orbital analysis

The highest occupied molecular orbitals (HOMOs) and the lowest unoccupied molecular orbitals (LUMOs) are named as frontier molecular orbitals (FMOs). The FMOs play an important role in the optical and electric properties. The frontier orbital gap characterizes the chemical reactivity and the kinetic stability of the molecule. A molecule with a small frontier orbital gap is generally associated with a high chemical reactivity, low kinetic stability and is also termed a soft molecule. The density functional theory (DFT) quantum-chemical calculations for the title compound were performed at the B3LYP/6–311 G(d,p) level (Becke, 1993) as implemented in *GAUSSIAN09* (Frisch *et al.*, 2009). Fig. 3 illustrates the HOMO and LUMO energy levels of the title compound. The small HOMO–LUMO energy gap (3.19 eV) in this compound indicates the chemical reactivity is strong and the kinetic stability is weak, which in turn increases the non-linear optical activity. As a result, with the small HOMO–LUMO energy gap, this compound could potentially be used in optoelectronic applications.

6. Molecular electrostatic potential (MEP)

The MEP map of the title molecule was calculated theoretically at the B3LYP/6–311G(d,p) level of theory and is illustrated in Fig. 4. The blue-coloured region is electrophilic

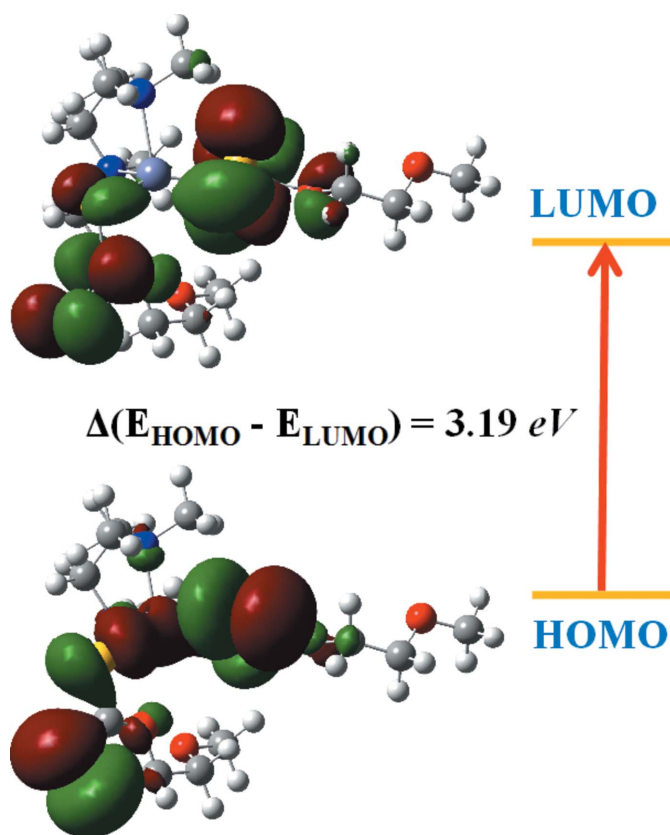


Figure 3
The electron distribution of the HOMO and LUMO energy levels of the title compound.

and electron poor, whereas the red colour indicates the nucleophilic region with rich electrons in the environment and provide information about interactions that can occur between molecules (Tankov & Yankova, 2019). In the title compound, the reactive sites are localized near the C–O group: this is the region having the most negative potential spots (red colour), all over the oxygen atom because of the

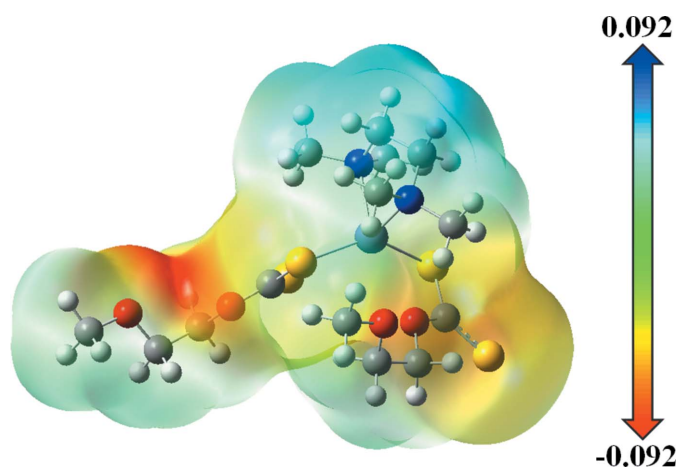


Figure 4
The total electron density three-dimensional surface mapped for the title compound with the electrostatic potential calculated at the B3LYP/6–311 G(d,p) level.

C—H...O interactions in the crystal structure. The negative potential value of -0.092 a.u. indicates the region that shows the strongest repulsion (electrophilic attack). In addition, the most positive region is located at the hydrogen atoms and shows the strongest attraction (nucleophilic attack) sites, which involve the N,N,N',N' -tetramethylethylenediamine moiety.

7. Synthesis and crystallization

Tetramethylethylenediamine (10 mmol, 1.16 g) was added to a hot solution of $Zn(CH_3CO_2)_2 \cdot 2H_2O$ (10 mmol, 2.20 g) in 2-methoxyethanol. A hot solution of potassium 2-methoxyethylxanthate (20 mmol, 3.81 g) in 2-methoxyethanol was added and the mixture was stirred for 30 min. Water was added to the mixture and a white precipitate was formed. The product was recrystallized from acetone.

8. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 3. The C-bound H atoms were positioned geometrically and refined using a riding model, with C—H = 0.98 and 0.99 Å and with $U_{iso}(H) = 1.5U_{eq}(C)$ for methyl H atoms and $1.2U_{eq}(C)$ otherwise. All atoms of the amine ligand are disordered and were modelled as two orientations with relative occupancies of 0.538 (6) and 0.462 (6). The diffuse electron density of half an acetone solvent molecule was removed with the solvent-mask procedure implemented in *OLEX2* (Dolomanov *et al.*, 2009). There are two voids of 122.4 \AA^3 in the unit cell and the electron count was 18.2 per void.

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Table 3

Experimental details.

Crystal data	
Chemical formula	$[Zn(C_4H_7O_2S_2)_2(C_6H_{16}N_2)] \cdot 0.5C_3H_6O$
M_r	513.05
Crystal system, space group	Monoclinic, $P2_1/n$
Temperature (K)	100
a, b, c (Å)	9.604 (3), 22.785 (6), 11.374 (3)
β (°)	106.304 (12)
V (Å ³)	2389.0 (12)
Z	4
Radiation type	Mo $K\alpha$
μ (mm ⁻¹)	1.40
Crystal size (mm)	$0.56 \times 0.52 \times 0.06$
Data collection	
Diffractometer	Bruker APEXII CCD
Absorption correction	Multi-scan (<i>SADABS</i> ; Bruker, 2009)
T_{min}, T_{max}	0.594, 0.746
No. of measured, independent and observed [$I > 2\sigma(I)$] reflections	35197, 5276, 3870
R_{int}	0.054
$(\sin \theta/\lambda)_{max}$ (Å ⁻¹)	0.650
Refinement	
$R[F^2 > 2\sigma(F^2)], wR(F^2), S$	0.040, 0.105, 1.06
No. of reflections	5276
No. of parameters	299
No. of restraints	244
H-atom treatment	H-atom parameters constrained
$\Delta\rho_{max}, \Delta\rho_{min}$ (e Å ⁻³)	0.41, -0.69

Computer programs: *APEX2* and *SAINT* (Bruker, 2009), *SHELXT* (Sheldrick, 2015a), *SHELXL* (Sheldrick, 2015b) and *OLEX2* (Dolomanov *et al.*, 2009).

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supporting information

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Computing details

Data collection: *APEX2* (Bruker, 2009); cell refinement: *SAINT* (Bruker, 2009); data reduction: *SAINT* (Bruker, 2009); program(s) used to solve structure: *SHELXT* (Sheldrick, 2015a); program(s) used to refine structure: *SHELXL* (Sheldrick, 2015b); molecular graphics: *OLEX2* (Dolomanov *et al.*, 2009); software used to prepare material for publication: *OLEX2* (Dolomanov *et al.*, 2009).

Bis(2-methoxyethyl xanthato- κ S)(*N,N,N',N'*-tetramethylethylenediamine- κ^2 *N,N'*)zinc(II) acetone hemisolvate

Crystal data

[Zn(C₄H₇O₂S₂)₂(C₆H₁₆N₂)]·0.5C₃H₆O

M_r = 513.05

Monoclinic, *P2₁/n*

a = 9.604 (3) Å

b = 22.785 (6) Å

c = 11.374 (3) Å

β = 106.304 (12)°

V = 2389.0 (12) Å³

Z = 4

F(000) = 1016

D_x = 1.346 Mg m⁻³

Mo *K* α radiation, λ = 0.71073 Å

Cell parameters from 8223 reflections

θ = 2.4–27.2°

μ = 1.40 mm⁻¹

T = 100 K

Plate, colourless

0.56 × 0.52 × 0.06 mm

Data collection

Bruker APEXII CCD
diffractometer

φ and ω scans

Absorption correction: multi-scan
(SADABS; Bruker, 2009)

T_{min} = 0.594, *T_{max}* = 0.746

35197 measured reflections

5276 independent reflections

3870 reflections with *I* > 2 σ (*I*)

R_{int} = 0.054

θ_{\max} = 27.5°, θ_{\min} = 2.6°

h = -12→12

k = -29→29

l = -13→14

Refinement

Refinement on *F*²

Least-squares matrix: full

R[*F*² > 2 σ (*F*²)] = 0.040

wR(*F*²) = 0.105

S = 1.06

5276 reflections

299 parameters

244 restraints

Hydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained

w = 1/[$\sigma^2(F_o^2) + (0.057P)^2$]

where *P* = (*F_o*² + 2*F_c*²)/3

(Δ/σ)_{max} = 0.001

$\Delta\rho_{\max}$ = 0.41 e Å⁻³

$\Delta\rho_{\min}$ = -0.68 e Å⁻³

Special details

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

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

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
Zn1	0.94679 (3)	0.20615 (2)	0.25390 (3)	0.01743 (10)	
S1	0.77332 (8)	0.16292 (3)	0.09443 (7)	0.02362 (17)	
S2	0.81448 (9)	0.08333 (3)	-0.10110 (7)	0.02709 (19)	
S3	1.11500 (8)	0.15676 (3)	0.40643 (6)	0.02205 (17)	
S4	0.82433 (9)	0.13279 (3)	0.44905 (8)	0.0312 (2)	
N1	1.0669 (9)	0.2630 (3)	0.1684 (7)	0.0187 (17)	0.538 (6)
N2	0.8649 (9)	0.2832 (2)	0.3157 (8)	0.0199 (19)	0.538 (6)
C1	1.0675 (9)	0.2455 (3)	0.0430 (7)	0.0234 (15)	0.538 (6)
H1A	0.9674	0.2412	-0.0084	0.035*	0.538 (6)
H1B	1.1167	0.2757	0.0081	0.035*	0.538 (6)
H1C	1.1187	0.2081	0.0465	0.035*	0.538 (6)
C2	1.2174 (11)	0.2697 (6)	0.2467 (12)	0.022 (2)	0.538 (6)
H2A	1.2162	0.2813	0.3294	0.034*	0.538 (6)
H2B	1.2689	0.2323	0.2505	0.034*	0.538 (6)
H2C	1.2669	0.3000	0.2121	0.034*	0.538 (6)
C3	0.9856 (7)	0.3192 (2)	0.1604 (6)	0.0229 (13)	0.538 (6)
H3A	0.8955	0.3170	0.0919	0.027*	0.538 (6)
H3B	1.0453	0.3517	0.1431	0.027*	0.538 (6)
C4	0.9477 (7)	0.3317 (2)	0.2780 (5)	0.0245 (13)	0.538 (6)
H4A	1.0384	0.3385	0.3442	0.029*	0.538 (6)
H4B	0.8894	0.3681	0.2679	0.029*	0.538 (6)
C5	0.7089 (10)	0.2896 (4)	0.2515 (12)	0.022 (2)	0.538 (6)
H5A	0.6950	0.2885	0.1628	0.033*	0.538 (6)
H5B	0.6548	0.2573	0.2750	0.033*	0.538 (6)
H5C	0.6736	0.3271	0.2740	0.033*	0.538 (6)
C6	0.883 (2)	0.2849 (6)	0.4502 (9)	0.029 (3)	0.538 (6)
H6A	0.9861	0.2808	0.4943	0.044*	0.538 (6)
H6B	0.8470	0.3225	0.4719	0.044*	0.538 (6)
H6C	0.8282	0.2527	0.4729	0.044*	0.538 (6)
N1A	1.0736 (10)	0.2669 (3)	0.1831 (8)	0.019 (2)	0.462 (6)
N2A	0.8391 (10)	0.2822 (3)	0.2930 (9)	0.020 (2)	0.462 (6)
C1A	1.0349 (10)	0.2672 (4)	0.0475 (7)	0.0209 (17)	0.462 (6)
H1AA	0.9303	0.2736	0.0142	0.031*	0.462 (6)
H1AB	1.0877	0.2988	0.0201	0.031*	0.462 (6)
H1AC	1.0611	0.2294	0.0184	0.031*	0.462 (6)
C2A	1.2323 (12)	0.2571 (7)	0.2278 (15)	0.026 (3)	0.462 (6)
H2AA	1.2624	0.2566	0.3175	0.039*	0.462 (6)
H2AB	1.2564	0.2194	0.1968	0.039*	0.462 (6)
H2AC	1.2830	0.2888	0.1985	0.039*	0.462 (6)

C3A	1.0409 (7)	0.3254 (3)	0.2283 (8)	0.0261 (15)	0.462 (6)
H3AA	1.0678	0.3568	0.1787	0.031*	0.462 (6)
H3AB	1.0991	0.3307	0.3145	0.031*	0.462 (6)
C4A	0.8818 (7)	0.3300 (3)	0.2199 (7)	0.0258 (15)	0.462 (6)
H4AA	0.8618	0.3687	0.2514	0.031*	0.462 (6)
H4AB	0.8237	0.3269	0.1332	0.031*	0.462 (6)
C5A	0.6793 (11)	0.2766 (5)	0.2581 (15)	0.022 (2)	0.462 (6)
H5AA	0.6435	0.2671	0.1708	0.033*	0.462 (6)
H5AB	0.6520	0.2452	0.3063	0.033*	0.462 (6)
H5AC	0.6365	0.3137	0.2741	0.033*	0.462 (6)
C6A	0.889 (2)	0.2966 (7)	0.4260 (10)	0.023 (3)	0.462 (6)
H6AA	0.9949	0.3006	0.4516	0.035*	0.462 (6)
H6AB	0.8448	0.3336	0.4409	0.035*	0.462 (6)
H6AC	0.8603	0.2651	0.4731	0.035*	0.462 (6)
C7	0.8778 (3)	0.11821 (10)	0.0299 (3)	0.0221 (6)	
O8	1.0160 (2)	0.11540 (7)	0.09989 (17)	0.0220 (4)	
C9	1.1153 (3)	0.07649 (11)	0.0618 (3)	0.0262 (7)	
H9A	1.1345	0.0914	-0.0139	0.031*	
H9B	1.0731	0.0367	0.0454	0.031*	
C10	1.2527 (3)	0.07470 (11)	0.1641 (3)	0.0246 (6)	
H10A	1.2301	0.0685	0.2429	0.030*	
H10B	1.3139	0.0417	0.1514	0.030*	
O11	1.3286 (2)	0.12830 (8)	0.1677 (2)	0.0318 (5)	
C12	1.4621 (3)	0.12837 (14)	0.2611 (3)	0.0380 (8)	
H12A	1.5113	0.1660	0.2607	0.057*	
H12B	1.4437	0.1226	0.3408	0.057*	
H12C	1.5237	0.0965	0.2463	0.057*	
C13	1.0016 (3)	0.12252 (11)	0.4785 (3)	0.0208 (6)	
O14	1.0774 (2)	0.08476 (8)	0.56361 (18)	0.0248 (4)	
C15	1.0045 (4)	0.05469 (13)	0.6430 (3)	0.0303 (7)	
H15A	0.9223	0.0311	0.5933	0.036*	
H15B	0.9664	0.0836	0.6911	0.036*	
C16	1.1136 (4)	0.01571 (12)	0.7265 (3)	0.0330 (8)	
H16A	1.0658	-0.0086	0.7762	0.040*	
H16B	1.1574	-0.0109	0.6779	0.040*	
O17	1.2226 (2)	0.05108 (8)	0.80437 (19)	0.0278 (5)	
C18	1.3393 (4)	0.01777 (13)	0.8787 (3)	0.0378 (8)	
H18A	1.4112	0.0443	0.9305	0.057*	
H18B	1.3026	-0.0089	0.9307	0.057*	
H18C	1.3847	-0.0050	0.8264	0.057*	

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Zn1	0.01874 (19)	0.01410 (15)	0.01842 (19)	0.00001 (12)	0.00353 (13)	-0.00176 (11)
S1	0.0204 (4)	0.0212 (3)	0.0258 (4)	0.0008 (3)	0.0008 (3)	-0.0072 (3)
S2	0.0331 (5)	0.0199 (3)	0.0239 (4)	-0.0004 (3)	0.0008 (3)	-0.0061 (3)
S3	0.0203 (4)	0.0247 (3)	0.0203 (4)	0.0006 (3)	0.0042 (3)	0.0036 (3)

S4	0.0221 (4)	0.0326 (4)	0.0391 (5)	-0.0011 (3)	0.0089 (4)	-0.0034 (3)
N1	0.019 (3)	0.014 (3)	0.024 (3)	0.004 (2)	0.006 (3)	0.002 (2)
N2	0.023 (3)	0.016 (2)	0.018 (4)	-0.002 (2)	0.002 (3)	-0.003 (2)
C1	0.029 (4)	0.022 (4)	0.023 (3)	0.001 (3)	0.012 (3)	-0.003 (3)
C2	0.023 (4)	0.022 (5)	0.024 (5)	0.001 (3)	0.009 (3)	0.001 (3)
C3	0.027 (3)	0.013 (2)	0.029 (3)	0.003 (2)	0.009 (2)	0.003 (2)
C4	0.028 (3)	0.016 (2)	0.030 (3)	-0.002 (2)	0.009 (3)	-0.005 (2)
C5	0.019 (4)	0.011 (4)	0.034 (4)	-0.006 (3)	0.006 (3)	-0.003 (3)
C6	0.045 (5)	0.028 (5)	0.014 (4)	0.002 (4)	0.008 (4)	-0.008 (4)
N1A	0.022 (3)	0.016 (3)	0.020 (3)	-0.001 (3)	0.004 (3)	-0.001 (3)
N2A	0.025 (4)	0.016 (3)	0.016 (4)	-0.002 (2)	0.002 (3)	-0.009 (2)
C1A	0.024 (4)	0.024 (4)	0.016 (3)	-0.001 (3)	0.008 (3)	-0.005 (3)
C2A	0.013 (4)	0.033 (7)	0.030 (6)	-0.004 (4)	0.001 (4)	0.002 (4)
C3A	0.034 (3)	0.015 (2)	0.031 (3)	-0.002 (2)	0.010 (3)	-0.005 (2)
C4A	0.033 (3)	0.017 (2)	0.025 (3)	0.005 (3)	0.004 (3)	-0.002 (2)
C5A	0.016 (5)	0.013 (5)	0.033 (5)	-0.004 (3)	0.000 (4)	-0.006 (3)
C6A	0.023 (4)	0.033 (6)	0.014 (5)	-0.001 (4)	0.003 (4)	-0.005 (4)
C7	0.0281 (17)	0.0105 (11)	0.0239 (16)	-0.0027 (11)	0.0008 (13)	0.0009 (10)
O8	0.0218 (11)	0.0173 (9)	0.0240 (11)	0.0031 (7)	0.0019 (9)	-0.0058 (7)
C9	0.0274 (17)	0.0213 (13)	0.0292 (17)	0.0043 (12)	0.0070 (13)	-0.0070 (11)
C10	0.0270 (17)	0.0133 (12)	0.0343 (18)	0.0000 (11)	0.0097 (13)	-0.0023 (11)
O11	0.0291 (13)	0.0238 (10)	0.0375 (13)	-0.0099 (9)	0.0012 (10)	0.0063 (9)
C12	0.0253 (19)	0.0333 (17)	0.049 (2)	-0.0047 (13)	-0.0001 (16)	-0.0026 (14)
C13	0.0247 (16)	0.0187 (12)	0.0174 (15)	-0.0010 (11)	0.0033 (12)	-0.0016 (10)
O14	0.0255 (12)	0.0254 (10)	0.0253 (12)	-0.0016 (8)	0.0100 (9)	0.0046 (8)
C15	0.037 (2)	0.0319 (15)	0.0236 (18)	-0.0143 (13)	0.0105 (14)	0.0024 (12)
C16	0.052 (2)	0.0203 (14)	0.0284 (18)	-0.0091 (14)	0.0133 (15)	0.0034 (12)
O17	0.0337 (13)	0.0198 (9)	0.0277 (12)	0.0002 (8)	0.0051 (10)	0.0041 (8)
C18	0.044 (2)	0.0318 (16)	0.037 (2)	0.0101 (14)	0.0098 (16)	0.0124 (14)

Geometric parameters (Å, °)

Zn1—S1	2.3107 (9)	C1A—H1AB	0.9800
Zn1—S3	2.3050 (9)	C1A—H1AC	0.9800
Zn1—N1	2.141 (5)	C2A—H2AA	0.9800
Zn1—N2	2.123 (5)	C2A—H2AB	0.9800
Zn1—N1A	2.144 (6)	C2A—H2AC	0.9800
Zn1—N2A	2.128 (6)	C3A—H3AA	0.9900
S1—C7	1.731 (3)	C3A—H3AB	0.9900
S2—C7	1.647 (3)	C3A—C4A	1.508 (7)
S3—C13	1.723 (3)	C4A—H4AA	0.9900
S4—C13	1.657 (3)	C4A—H4AB	0.9900
N1—C1	1.482 (6)	C5A—H5AA	0.9800
N1—C2	1.477 (7)	C5A—H5AB	0.9800
N1—C3	1.489 (6)	C5A—H5AC	0.9800
N2—C4	1.492 (6)	C6A—H6AA	0.9800
N2—C5	1.478 (7)	C6A—H6AB	0.9800
N2—C6	1.491 (7)	C6A—H6AC	0.9800

C1—H1A	0.9800	C7—O8	1.344 (3)
C1—H1B	0.9800	O8—C9	1.454 (3)
C1—H1C	0.9800	C9—H9A	0.9900
C2—H2A	0.9800	C9—H9B	0.9900
C2—H2B	0.9800	C9—C10	1.495 (4)
C2—H2C	0.9800	C10—H10A	0.9900
C3—H3A	0.9900	C10—H10B	0.9900
C3—H3B	0.9900	C10—O11	1.417 (3)
C3—C4	1.509 (6)	O11—C12	1.417 (4)
C4—H4A	0.9900	C12—H12A	0.9800
C4—H4B	0.9900	C12—H12B	0.9800
C5—H5A	0.9800	C12—H12C	0.9800
C5—H5B	0.9800	C13—O14	1.346 (3)
C5—H5C	0.9800	O14—C15	1.459 (3)
C6—H6A	0.9800	C15—H15A	0.9900
C6—H6B	0.9800	C15—H15B	0.9900
C6—H6C	0.9800	C15—C16	1.494 (4)
N1A—C1A	1.482 (7)	C16—H16A	0.9900
N1A—C2A	1.482 (7)	C16—H16B	0.9900
N1A—C3A	1.494 (7)	C16—O17	1.418 (4)
N2A—C4A	1.496 (7)	O17—C18	1.420 (4)
N2A—C5A	1.479 (7)	C18—H18A	0.9800
N2A—C6A	1.490 (7)	C18—H18B	0.9800
C1A—H1AA	0.9800	C18—H18C	0.9800
S3—Zn1—S1	125.54 (3)	H1AB—C1A—H1AC	109.5
N1—Zn1—S1	105.2 (2)	N1A—C2A—H2AA	109.5
N1—Zn1—S3	106.5 (2)	N1A—C2A—H2AB	109.5
N2—Zn1—S1	111.1 (2)	N1A—C2A—H2AC	109.5
N2—Zn1—S3	113.7 (2)	H2AA—C2A—H2AB	109.5
N2—Zn1—N1	86.9 (2)	H2AA—C2A—H2AC	109.5
N1A—Zn1—S1	109.9 (3)	H2AB—C2A—H2AC	109.5
N1A—Zn1—S3	104.2 (3)	N1A—C3A—H3AA	109.6
N2A—Zn1—S1	103.0 (3)	N1A—C3A—H3AB	109.6
N2A—Zn1—S3	121.3 (3)	N1A—C3A—C4A	110.4 (6)
N2A—Zn1—N1A	85.1 (3)	H3AA—C3A—H3AB	108.1
C7—S1—Zn1	101.98 (10)	C4A—C3A—H3AA	109.6
C13—S3—Zn1	100.13 (10)	C4A—C3A—H3AB	109.6
C1—N1—Zn1	115.0 (5)	N2A—C4A—C3A	110.2 (6)
C1—N1—C3	108.4 (5)	N2A—C4A—H4AA	109.6
C2—N1—Zn1	110.4 (8)	N2A—C4A—H4AB	109.6
C2—N1—C1	109.9 (7)	C3A—C4A—H4AA	109.6
C2—N1—C3	110.8 (7)	C3A—C4A—H4AB	109.6
C3—N1—Zn1	102.0 (4)	H4AA—C4A—H4AB	108.1
C4—N2—Zn1	103.9 (4)	N2A—C5A—H5AA	109.5
C5—N2—Zn1	109.6 (7)	N2A—C5A—H5AB	109.5
C5—N2—C4	109.5 (6)	N2A—C5A—H5AC	109.5
C5—N2—C6	108.4 (9)	H5AA—C5A—H5AB	109.5

C6—N2—Zn1	114.5 (7)	H5AA—C5A—H5AC	109.5
C6—N2—C4	110.9 (7)	H5AB—C5A—H5AC	109.5
N1—C1—H1A	109.5	N2A—C6A—H6AA	109.5
N1—C1—H1B	109.5	N2A—C6A—H6AB	109.5
N1—C1—H1C	109.5	N2A—C6A—H6AC	109.5
H1A—C1—H1B	109.5	H6AA—C6A—H6AB	109.5
H1A—C1—H1C	109.5	H6AA—C6A—H6AC	109.5
H1B—C1—H1C	109.5	H6AB—C6A—H6AC	109.5
N1—C2—H2A	109.5	S2—C7—S1	123.82 (18)
N1—C2—H2B	109.5	O8—C7—S1	111.76 (19)
N1—C2—H2C	109.5	O8—C7—S2	124.4 (2)
H2A—C2—H2B	109.5	C7—O8—C9	118.3 (2)
H2A—C2—H2C	109.5	O8—C9—H9A	110.3
H2B—C2—H2C	109.5	O8—C9—H9B	110.3
N1—C3—H3A	109.4	O8—C9—C10	107.1 (2)
N1—C3—H3B	109.4	H9A—C9—H9B	108.5
N1—C3—C4	111.1 (5)	C10—C9—H9A	110.3
H3A—C3—H3B	108.0	C10—C9—H9B	110.3
C4—C3—H3A	109.4	C9—C10—H10A	109.8
C4—C3—H3B	109.4	C9—C10—H10B	109.8
N2—C4—C3	113.3 (5)	H10A—C10—H10B	108.2
N2—C4—H4A	108.9	O11—C10—C9	109.5 (2)
N2—C4—H4B	108.9	O11—C10—H10A	109.8
C3—C4—H4A	108.9	O11—C10—H10B	109.8
C3—C4—H4B	108.9	C12—O11—C10	111.9 (2)
H4A—C4—H4B	107.7	O11—C12—H12A	109.5
N2—C5—H5A	109.5	O11—C12—H12B	109.5
N2—C5—H5B	109.5	O11—C12—H12C	109.5
N2—C5—H5C	109.5	H12A—C12—H12B	109.5
H5A—C5—H5B	109.5	H12A—C12—H12C	109.5
H5A—C5—H5C	109.5	H12B—C12—H12C	109.5
H5B—C5—H5C	109.5	S4—C13—S3	126.24 (16)
N2—C6—H6A	109.5	O14—C13—S3	110.2 (2)
N2—C6—H6B	109.5	O14—C13—S4	123.6 (2)
N2—C6—H6C	109.5	C13—O14—C15	119.2 (2)
H6A—C6—H6B	109.5	O14—C15—H15A	110.2
H6A—C6—H6C	109.5	O14—C15—H15B	110.2
H6B—C6—H6C	109.5	O14—C15—C16	107.5 (3)
C1A—N1A—Zn1	113.0 (6)	H15A—C15—H15B	108.5
C1A—N1A—C3A	109.8 (6)	C16—C15—H15A	110.2
C2A—N1A—Zn1	114.3 (9)	C16—C15—H15B	110.2
C2A—N1A—C1A	106.9 (8)	C15—C16—H16A	109.9
C2A—N1A—C3A	108.2 (7)	C15—C16—H16B	109.9
C3A—N1A—Zn1	104.5 (4)	H16A—C16—H16B	108.3
C4A—N2A—Zn1	104.6 (4)	O17—C16—C15	108.8 (2)
C5A—N2A—Zn1	113.7 (7)	O17—C16—H16A	109.9
C5A—N2A—C4A	109.8 (7)	O17—C16—H16B	109.9
C5A—N2A—C6A	107.8 (9)	C16—O17—C18	112.9 (2)

C6A—N2A—Zn1	110.8 (9)	O17—C18—H18A	109.5
C6A—N2A—C4A	110.1 (8)	O17—C18—H18B	109.5
N1A—C1A—H1AA	109.5	O17—C18—H18C	109.5
N1A—C1A—H1AB	109.5	H18A—C18—H18B	109.5
N1A—C1A—H1AC	109.5	H18A—C18—H18C	109.5
H1AA—C1A—H1AB	109.5	H18B—C18—H18C	109.5
H1AA—C1A—H1AC	109.5		
Zn1—S1—C7—S2	-170.71 (15)	C2—N1—C3—C4	-73.8 (9)
Zn1—S1—C7—O8	9.49 (19)	C5—N2—C4—C3	-84.1 (8)
Zn1—S3—C13—S4	-6.90 (19)	C6—N2—C4—C3	156.4 (9)
Zn1—S3—C13—O14	172.51 (16)	N1A—C3A—C4A—N2A	58.4 (9)
Zn1—N1—C3—C4	43.8 (6)	C1A—N1A—C3A—C4A	81.0 (8)
Zn1—N2—C4—C3	32.9 (7)	C2A—N1A—C3A—C4A	-162.6 (10)
Zn1—N1A—C3A—C4A	-40.5 (8)	C5A—N2A—C4A—C3A	-164.5 (9)
Zn1—N2A—C4A—C3A	-42.1 (8)	C6A—N2A—C4A—C3A	77.0 (10)
S1—C7—O8—C9	176.11 (18)	C7—O8—C9—C10	-171.4 (2)
S2—C7—O8—C9	-3.7 (3)	O8—C9—C10—O11	-73.4 (3)
S3—C13—O14—C15	174.79 (18)	C9—C10—O11—C12	-178.4 (2)
S4—C13—O14—C15	-5.8 (3)	C13—O14—C15—C16	179.5 (2)
N1—C3—C4—N2	-55.6 (8)	O14—C15—C16—O17	65.4 (3)
C1—N1—C3—C4	165.5 (6)	C15—C16—O17—C18	-173.9 (3)

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>
C1—H1C...O8	0.98	2.48	3.103 (7)	121
C2A—H2AB...O11	0.98	2.24	3.207 (13)	168
C5A—H5AA...S1	0.98	2.92	3.454 (16)	115
C6—H6C...S4	0.98	2.74	3.512 (13)	136
C6—H6B...O11 ⁱ	0.98	2.54	3.321 (13)	136
C3A—H3AB...S2 ⁱⁱ	0.99	2.81	3.483 (7)	125
C6A—H6AA...S1 ⁱⁱ	0.98	2.84	3.764 (16)	158
C4A—H4AA...O17 ⁱⁱⁱ	0.99	2.44	3.380 (6)	159
C4A—H4AB...S3 ⁱⁱⁱ	0.99	2.81	3.774 (8)	164
C9—H9A...O17 ^{iv}	0.99	2.61	3.415 (4)	138
C9—H9B...S2 ^v	0.99	2.94	3.708 (3)	135
C18—H18B...S2 ^{vi}	0.98	3.02	3.998 (3)	176

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