Acta Crystallographica Section E **Structure Reports** Online

ISSN 1600-5368

Bis(isopropylammonium) tetrasulfidomolybdate(VI)

Bikshandarkoil R. Srinivasan,^a* Christian Näther,^b Ashish R. Naik^a and Wolfgang Bensch^b

^aDepartment of Chemistry, Goa University, Goa 403206, India, and ^bInstitut für Anorganische Chemie, Christian-Albrechts-Universität Kiel, Olshausenstr, 40. D-24098 Kiel, Germany Correspondence e-mail: srini@unigoa.ac.in

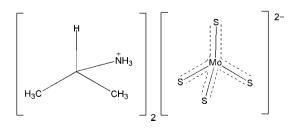
Received 22 November 2007; accepted 26 November 2007

Key indicators: single-crystal X-ray study; T = 170 K; mean σ (C–C) = 0.004 Å; R factor = 0.026; wR factor = 0.071; data-to-parameter ratio = 25.1.

The title compound, $(C_3H_{10}N)_2[MoS_4]$, was synthesized by passing a rapid stream of H₂S into an aqueous isopropylamine solution of molybdic acid. The title compound is isotypic with the corresponding W analogue $(C_3H_{10}N)_2[WS_4]$; its structure consists of a slightly distorted tetrahedral $[MoS_4]^{2-}$ dianion crystallographically independent isopropyland two ammonium cations, with all atoms located in general positions. The cations and anion are linked by weak N-H···S and C- $H \cdots S$ interactions, the strength and number of which can explain the observed Mo-S bond distances.

Related literature

Previous reports give details of the structural characterization of several organic ammonium tetrasulfidomolybdates derived from chiral amines (Srinivasan, Naik et al., 2007), diamines (Srinivasan et al., 2001; Srinivasan, Dhuri et al., 2005; Srinivasan, Näther & Bensch 2005), triamines (Srinivasan, Dhuri et al., 2007), cyclic amines (Srinivasan, Näther & Bensch 2006), a tetraamine (Srinivasan et al., 2004), a primary amine (Srinivasan, Näther, Naik & Bensch 2006) and a secondary amine (Srinivasan, Girkar & Raghavaiah 2007). The title compound analogue isotypic with the corresponding W is (C₃H₁₀N)₂[WS₄] (Srinivasan, Näther, Dhuri & Bensch 2006).



Experimental

Crystal data

$(C_{3}H_{10}N)_{2}[MoS_{4}]$	V = 2937.3 (4) Å ³
$M_r = 344.42$	Z = 8
Monoclinic, $C2/c$	Mo $K\alpha$ radiation
a = 20.2640 (14) Å	$\mu = 1.43 \text{ mm}^{-1}$
b = 13.9118 (12) Å	T = 170 (2) K
c = 11.0933 (8) Å	$0.13 \times 0.1 \times 0.08 \text{ mm}$
$\beta = 110.076 \ (8)^{\circ}$	

Data collection

Stoe IPDS 1 diffractometer Absorption correction: numerical (X-SHAPE; Stoe & Cie, 1998) $T_{\min} = 0.756, T_{\max} = 0.830$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.026$	125 parameters
$wR(F^2) = 0.071$	H-atom parameters constrained
S = 1.03	$\Delta \rho_{\rm max} = 0.64 \text{ e } \text{\AA}^{-3}$
3134 reflections	$\Delta \rho_{\rm min} = -0.72 \ {\rm e} \ {\rm \AA}^{-3}$

10806 measured reflections

 $R_{\rm int} = 0.025$

3134 independent reflections

2716 reflections with $I > 2\sigma(I)$

Table 1

Selected geometric parameters (Å, °).

Mo-S2	2.1695 (6)	Mo-S1	2.2023 (6)
Mo-S3	2.1769 (5)	Mo-S4	2.2085 (6)
S2-Mo-S3	109.08 (2)	S2-Mo-S4	109.01 (2)
S2-Mo-S1	109.02 (3)	S3-Mo-S4	109.78 (2)
S3-Mo-S1	109.51 (2)	S1-Mo-S4	110.42 (2)

Table 2			
Hydrogen-bond	geometry	(Å,	°).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots \mathbf{A}$
$N1 - H1N1 \cdot \cdot \cdot S4$	0.91	2.43	3.310 (2)	162
$N1 - H2N1 \cdot \cdot \cdot S4^{i}$	0.91	2.47	3.359 (2)	165
$N1 - H3N1 \cdot \cdot \cdot S1^{ii}$	0.91	2.54	3.4297 (19)	165
$N1 - H3N1 \cdot \cdot \cdot S3^{ii}$	0.91	2.85	3.3019 (19)	112
$N2-H1N2\cdots S3^{iii}$	0.91	2.58	3.376 (2)	147
$N2-H1N2 \cdot \cdot \cdot S2^{iii}$	0.91	2.92	3.580 (2)	131
$N2-H2N2\cdots S4$	0.91	2.43	3.309 (2)	164
$N2-H3N2\cdots S1^{i}$	0.91	2.50	3.393 (2)	169
$C2-H2A\cdots S4^{i}$	0.98	2.97	3.759 (3)	139
$C5-H5B\cdots S3^{ii}$	0.98	2.98	3.639 (6)	126
Symmetry codes:	(i) $x, -y +$	$-1, z - \frac{1}{2};$ (ii) $-x + \frac{1}{2}, -y + \frac{3}{2},$	-z + 1; (iii)

 $-x + \frac{1}{2}, y - \frac{1}{2}, -z + \frac{3}{2}$

Data collection: IPDS Program Package (Stoe & Cie, 1998); cell refinement: IPDS Program Package; data reduction: IPDS Program Package; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: DIAMOND (Brandenburg, 1999); software used to prepare material for publication: CIFTAB in SHELXTL (Bruker, 1998).

This work was supported by the Department of Science and Technology (DST), New Delhi, under grant No. SR/S1/IC-41/ 2003. ARN thanks the Deutscher Akademisher Austauschdienst (DAAD), Bonn, for a short-term visit to the University of Kiel.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: SI2062).

References

- Brandenburg, K. (1999). *DIAMOND*. Release 2.1c. Crystal Impact GbR, Bonn, Germany.
- Bruker (1998). SHELXTL. Version 5.10. Bruker AXS Inc., Madison, Wisconsin, USA.
- Sheldrick, G. M. (1997). SHELXS97 and SHELXL97. University of Göttingen, Germany.
- Srinivasan, B. R., Dhuri, S. N., Naik, A. R., Näther, C. & Bensch, W. (2007). *Polyhedron*, 26. In the press. doi:10.1016/jj.poly.2007.08.023.
- Srinivasan, B. R., Dhuri, S. N., Näther, C. & Bensch, W. (2005). Inorg. Chim. Acta, 358, 279–287.

- Srinivasan, B. R., Dhuri, S. N., Poisot, M., Näther, C. & Bensch, W. (2004). Z. Naturforsch. Teil B, 59, 1083–1092.
- Srinivasan, B. R., Girkar, S. V. & Raghavaiah, P. (2007). Acta Cryst. E63, m2737–m2738.
- Srinivasan, B. R., Naik, A. R., Näther, C. & Bensch, W. (2007). Z. Anorg. Allg. Chem. 633, 582–588.
- Srinivasan, B. R., Näther, C. & Bensch, W. (2005). Acta Cryst. E61, m2454– m2456.
- Srinivasan, B. R., Näther, C. & Bensch, W. (2006). Acta Cryst. C62, m98-m101. Srinivasan, B. R., Näther, C., Dhuri, S. N. & Bensch, W. (2006). Monatsh.
- Chem. 137, 397–411.
- Srinivasan, B. R., N\u00e4ther, C., Naik, A. R. & Bensch, W. (2006). Acta Cryst. E62, m1635–m1637.
- Srinivasan, B. R., Vernekar, B. K. & Nagarajan, K. (2001). Indian J. Chem. Sect. A, 40, 563–567.
- Stoe & Cie (1998). X-SHAPE (Version 1.03) and IPDS Program Package. (Version 2.89). Stoe & Cie, Darmstadt, Germany.

Acta Cryst. (2008). E64, m66-m67 [doi:10.1107/S1600536807063337]

Bis(isopropylammonium) tetrasulfidomolybdate(VI)

B. R. Srinivasan, C. Näther, A. R. Naik and W. Bensch

Comment

As part of an ongoing research programme, we are investigating the synthesis and structural aspects of organic ammonium tetrasulfidometalates of the group VI metals Mo and W (Srinivasan, Naik *et al.*, 2007). In earlier work we have structurally characterized several $[MoS_4]^{2-}$ compounds linked to organic cations derived from chiral amines (Srinivasan, Naik *et al.*, 2007), diamines (Srinivasan *et al.*, 2001; Srinivasan, Dhuri *et al.*, 2005; Srinivasan, Näther & Bensch 2005), triamines (Srinivasan, Dhuri *et al.*, 2007), cyclic amines (Srinivasan, Näther & Bensch 2006), a tetraamine (Srinivasan *et al.*, 2004), a primary amine (Srinivasan, Näther, Naik & Bensch 2006) and a secondary amine (Srinivasan, Girkar & Raghavaiah 2007). All the organic ammonium tetrasulfidomolybdates exhibit several weak hydrogen bonding interactions between the organic cations and $[MoS_4]^{2-}$ anions. We have shown that in some organic $[MoS_4]^{2-}$ compounds the organic amines are partially protonated (Srinivasan, Dhuri *et al.*, 2007). In the present work, we have employed isopropylamine (ipNH₂) for the synthesis of the title compound, which is isostructural with the corresponding W compound (C₃H₁₀N)₂[WS₄] (Srinivasan, Näther, Dhuri & Bensch 2006).

The structure of the title compound consists of discrete tetrahedral $[MoS_4]^{2-}$ ions and two crystallographically independent isopropylammonium cations (ipNH₂)⁺ (Fig. 1), with all atoms located in general positions. The geometric parameters of the organic cations are in agreement with those reported for the analogous $[WS_4]^{2-}$ compound. The MoS₄ tetrahedron is slightly distorted with S-Mo-S angles between 109.01 (2) and 110.42 (2) ° (Table 1). The Mo-S bond distances range from 2.1695 (6) to 2.2085 (6) Å with an average value of 2.1893 Å and are comparable to the bond lengths observed in several reported tetrathiomolybdates (Srinivasan, Dhuri et al., 2007). Two of the Mo-S bond lengths are shorter than the average Mo-S distance while the other two are longer. The weak H-bonding interactions between the cations and anions can explain the observed short and long Mo-S bond distances. A scrutiny of the structure reveals that the organic cations and tetrathiomolybdate anions are linked with the aid of several N-H···S and C-H···S hydrogen bonding interactions (Table 2). Thus each $[MoS_4]^{2-}$ is hydrogen bonded to seven different organic cations with the aid of eight N—H...S bonds and two weak C-H...S interactions (Fig.2). An examination of the surroundings of the cations reveals that one organic cation (N1) is H-bonded to three different $[MoS_4]^{2-}$ ions while the second organic cation (N2) is surrounded by four different $[MoS_4]^{2-}$ ions (Table 2). One H atom on each N atom functions as a bifurcated donor with the other two functioning as singly shared donors. One H atom attached to a methyl group from each unique cation is involved in a weak C—H···S interaction. S4 atom which makes the longest Mo-S bond at 2.2085 (6) Å is involved in three singly shared N-H···S bonds. S4 also makes the shortest singly shared N—H...S bond at 2.43 Å, which can explain the elongation of this bond. In contrast, S2 atom involved in the shortest Mo—S bond at 2.1695 (6) Å makes a bifurcated N—H…S bond at a longer S…H distance accompanied by a small NH-S angle. As a result of the H-bonding interactions, the cations and anions are organized such that the organic ammonium ions always point towards the S atoms of $[MoS_4]^{2-}$ as is evident in the observed sequence \cdots (ipNH₂)⁺ \cdots [MoS₄]²⁻ \cdots (ipNH₂)⁺ (ipNH₂)⁺ \cdots [MoS₄]²⁻ \cdots (ipNH₂)⁺ and so on when viewed along the *c* axis (Fig. 3). The

observed difference Δ between the longest and the shortest Mo—S bond of 0.0390 Å in the title compound is quite longer than the Δ value of 0.0334 Å in the isostructural tetrathiotungstate compound (C₃H₁₀N)₂[WS₄] (Srinivasan *et al.*, 2007*a*).

Experimental

A rapid stream of H_2S gas was passed into a solution containing molybdic acid (3 g) dissolved in water (30 ml) and isopropyl amine (10 ml). After 30 min when crystals begin to appear, the gas passing was stopped and the deep red reaction mixture was filtered. The filtrate was left aside for crystallization. After 3 to 4 h the crystalline product was filtered, washed with a little ice-cold water (2 ml), followed by 2-propanol (20 ml) and diethyl ether (10 ml) and dried to obtain 3.2 g of the title compound.

Refinement

All H atoms were located in difference map but were positioned with idealized geometry ((CH₃ and NH₃ allowed to rotate but not to tip) with 1.00 Å (C—H), 0.98 Å (methyl) and N—H = 0.91 Å) and were refined using a riding model, with U_{iso} (H) fixed at 1.5 U_{eq} (CH₃ and NH₃) and 1.5 U_{eq} (NH₃).

Figures

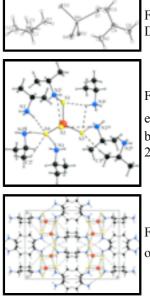


Fig. 1. The molecular structure of the title compound showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 50% probability level.

Fig. 2. A view of the surroundings of the $[MoS_4]^{2-}$ anion showing its linking to seven different (ipNH₂)⁺ cations with the aid of eight N—H…S and two C—H…S interactions, shown as broken lines. Symmetry codes: (i) x, -y + 1, z - 1/2 (ii) -x + 1/2, -y + 3/2, -z + 1; (iii) -x + 1/2, y - 1/2, -z + 3/2;

Fig. 3. A view along *c* axis of the crystallographic packing of the title compound showing the organization of the organic cations and $[MoS_4]^{2-}$ anions. H-bonds are shown as broken lines.

Bis(isopropylammonium) tetrasulfidomolybdate(VI)

Crystal data	
$(C_{3}H_{10}N)_{2}[MoS_{4}]$	$F_{000} = 1408$
$M_r = 344.42$	$D_{\rm x} = 1.558 { m Mg m}^{-3}$
Monoclinic, C2/c	Mo $K\alpha$ radiation $\lambda = 0.71073$ Å

Hall symbol: -C 2yc a = 20.2640 (14) Å b = 13.9118 (12) Å c = 11.0933 (8) Å $\beta = 110.076 (8)^{\circ}$ $V = 2937.3 (4) \text{ Å}^{3}$ Z = 8

Data collection

STOE IPDS 1 diffractometer	3134 independent reflections
Radiation source: fine-focus sealed tube	2716 reflections with $I > 2\sigma(I)$
Monochromator: graphite	$R_{\rm int} = 0.025$
T = 170(2) K	$\theta_{max} = 27.1^{\circ}$
Phi Scan scans	$\theta_{\min} = 2.4^{\circ}$
Absorption correction: numerical (X-SHAPE; Stoe & Cie, 1998)	$h = -23 \rightarrow 25$
$T_{\min} = 0.756, T_{\max} = 0.830$	$k = -17 \rightarrow 17$
10806 measured reflections	$l = -13 \rightarrow 14$

Refinement

Refinement on F^2	Hydrogen site location: inferred from neighbouring sites
Least-squares matrix: full	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.026$	$w = 1/[\sigma^2(F_0^2) + (0.0498P)^2 + 0.8301P]$ where $P = (F_0^2 + 2F_c^2)/3$
$wR(F^2) = 0.071$	$(\Delta/\sigma)_{\text{max}} = 0.001$
<i>S</i> = 1.03	$\Delta \rho_{max} = 0.64 \text{ e } \text{\AA}^{-3}$
3134 reflections	$\Delta \rho_{min} = -0.72 \text{ e} \text{ Å}^{-3}$
125 parameters	Extinction correction: SHELXL, $Fc^*=kFc[1+0.001xFc^2\lambda^3/sin(2\theta)]^{-1/4}$
Primary atom site location: structure-invariant direct methods	Extinction coefficient: 0.00132 (18)

Secondary atom site location: difference Fourier map

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.

Cell parameters from 8000 reflections

 $\theta = 12.5 - 28.0^{\circ}$

 $\mu = 1.43 \text{ mm}^{-1}$

T = 170 (2) K

 $0.13 \times 0.1 \times 0.08 \text{ mm}$

Needle, red

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 $F^2 > 2$ 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.

	x	у	Ζ	Uiso*/Ueq
Мо	0.277534 (8)	0.660872 (11)	0.763180 (16)	0.01395 (9)
S1	0.16255 (3)	0.66070 (4)	0.71330 (6)	0.02148 (14)
S2	0.32646 (3)	0.64606 (4)	0.96951 (5)	0.02172 (14)
S3	0.31057 (3)	0.79621 (4)	0.70333 (5)	0.02302 (14)
S4	0.31079 (3)	0.53985 (4)	0.66851 (6)	0.02458 (14)
N1	0.34948 (9)	0.61362 (13)	0.41696 (19)	0.0219 (4)
H1N1	0.3314	0.6038	0.4805	0.033*
H2N1	0.3318	0.5690	0.3541	0.033*
H3N1	0.3376	0.6735	0.3835	0.033*
C1	0.42792 (12)	0.60477 (18)	0.4708 (3)	0.0292 (6)
H1	0.4404	0.5389	0.5075	0.035*
C2	0.45832 (14)	0.61919 (19)	0.3658 (3)	0.0346 (6)
H2A	0.4421	0.5676	0.3023	0.052*
H2B	0.5097	0.6180	0.4027	0.052*
H2C	0.4429	0.6814	0.3241	0.052*
C3	0.45457 (18)	0.6778 (3)	0.5776 (3)	0.0557 (10)
H3A	0.4450	0.7428	0.5417	0.084*
H3B	0.5053	0.6695	0.6204	0.084*
H3C	0.4306	0.6684	0.6400	0.084*
N2	0.17411 (10)	0.40310 (14)	0.51513 (19)	0.0247 (4)
H1N2	0.1828	0.3534	0.5719	0.037*
H2N2	0.2062	0.4506	0.5478	0.037*
H3N2	0.1773	0.3820	0.4397	0.037*
C4	0.10154 (11)	0.44146 (15)	0.4922 (2)	0.0197 (4)
H4	0.0987	0.4646	0.5756	0.024*
C5	0.08879 (15)	0.52564 (17)	0.4005 (3)	0.0323 (6)
H5A	0.0933	0.5043	0.3195	0.048*
H5B	0.1234	0.5761	0.4386	0.048*
H5C	0.0414	0.5510	0.3842	0.048*
C6	0.04982 (13)	0.35982 (18)	0.4430 (3)	0.0307 (6)
H6A	0.0022	0.3828	0.4303	0.046*
H6B	0.0616	0.3073	0.5057	0.046*
H6C	0.0522	0.3366	0.3612	0.046*

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\hat{A}^2)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Мо	0.01492 (12)	0.01177 (12)	0.01591 (12)	-0.00135 (6)	0.00626 (8)	0.00189 (6)
S1	0.0147 (3)	0.0242 (3)	0.0243 (3)	-0.00166 (18)	0.0051 (2)	0.0059 (2)
S2	0.0215 (3)	0.0247 (3)	0.0167 (3)	-0.0033 (2)	0.0038 (2)	0.00321 (19)
S3	0.0266 (3)	0.0171 (3)	0.0275 (3)	-0.0053 (2)	0.0121 (2)	0.0054 (2)
S4	0.0337 (3)	0.0184 (3)	0.0257 (3)	0.0004 (2)	0.0154 (3)	-0.0027 (2)
N1	0.0182 (9)	0.0205 (9)	0.0258 (10)	-0.0003 (7)	0.0060 (8)	0.0033 (7)
C1	0.0181 (11)	0.0290 (12)	0.0371 (14)	0.0021 (9)	0.0049 (10)	0.0151 (10)

C2	0.0260 (12)	0.0357 (14)	0.0461 (16)	-0.0012 (10)	0.0174 (12)	-0.0046 (12)
C3	0.0378 (17)	0.103 (3)	0.0258 (14)	-0.0309 (17)	0.0098 (13)	-0.0058 (16)
N2	0.0184 (9)	0.0315 (10)	0.0242 (10)	-0.0023 (8)	0.0073 (8)	0.0044 (8)
C4	0.0190 (10)	0.0219 (10)	0.0211 (10)	-0.0015 (8)	0.0107 (9)	-0.0011 (8)
C5	0.0462 (15)	0.0203 (11)	0.0348 (13)	0.0051 (10)	0.0195 (12)	0.0037 (10)
C6	0.0205 (11)	0.0278 (12)	0.0444 (15)	-0.0041 (9)	0.0118 (11)	0.0000 (11)
Geometric	parameters (Å, °)					
Mo—S2		2.1695 (6)	C3—1	U2D	0.98	200
Mo—S2 Mo—S3		2.1769 (5)	C3—1		0.98	
Mo—S1		2.2023 (6)	N2—)2 (3)
Mo—S4		2.2025 (6)	N2—1		0.91	
N1-C1		1.499 (3)	N2—1		0.91	
NI—HINI		0.9100		H3N2	0.91	
N1—H2N1		0.9100	C4—(14 (3)
N1—H3N1		0.9100	C4—(14 (3)
C1-C2		1.506 (4)	C4—1		1.0	. ,
C1—C3		1.513 (4)	C5—1		0.98	
C1—H1		1.0000	C5—1		0.98	
C2—H2A		0.9800	C5—1		0.98	
C2—H2B		0.9800	C6—1		0.98	
C2—H2C		0.9800	C6—1		0.9800	
C3—H3A		0.9800	C6—1		0.9800	
S2—Mo—S	3	109.08 (2)		С3—НЗС	109	
S2—Mo—S S2—Mo—S		109.02 (3)		-C3-H3C	109	
S2—Mo—S S3—Mo—S		109.51 (2)			109	
S2—Mo—S		109.01 (2)		N2—H1N2	109	
S2 Mo S S3—Mo—S		109.78 (2)		N2—H2N2	109	
SJ Mo S S1—Mo—S		110.42 (2)		H1N2—N2—H2N2		
C1—N1—H		109.5		H1N2—N2—H2N2 109.5 C4—N2—H3N2 109.5		
C1—N1—H		109.5		H1N2—N2—H3N2 109.5		
H1N1—N1-		109.5		2—N2—H3N2	109	
C1—N1—H		109.5		C4—C5		.67 (19)
H1N1—N1-		109.5		C4—C6		.12 (18)
H2N1—N1-	—H3N1	109.5		C4—C6		.6 (2)
N1—C1—C	22	109.9 (2)	N2—	С4—Н4	108	
N1—C1—C	23	107.5 (2)	C5—4	С4—Н4	108	.8
C2—C1—C	3	112.7 (2)	C6—4	С4—Н4	108	.8
N1—C1—H	[1	108.9	C4—4	С5—Н5А	109	.5
С2—С1—Н	[1	108.9	C4—4	С5—Н5В	109	.5
С3—С1—Н	1	108.9	H5A-	—С5—Н5В	109	.5
С1—С2—Н	12A	109.5	C4—	С5—Н5С	109	.5
С1—С2—Н	2B	109.5	H5A-	—С5—Н5С	109	.5
H2A—C2—	-H2B	109.5	H5B-	—С5—Н5С	109	.5
С1—С2—Н	12C	109.5	C4—0	С6—Н6А	109	.5
H2A—C2—	-H2C	109.5	C4—0	С6—Н6В	109	.5
H2B—C2—	-H2C	109.5	H6A-	—С6—Н6В	109	.5
С1—С3—Н	[3A	109.5	C4—4	С6—Н6С	109	.5

C1—C3—H3B H3A—C3—H3B	109.5 109.5		Н6А—С6—Н6С Н6В—С6—Н6С		109.5 109.5	
Hydrogen-bond geometry (Å, °)						
D—H···A		<i>D</i> —Н	$H \cdots A$	$D \cdots A$	D—H··· A	
N1—H1N1…S4		0.91	2.43	3.310 (2)	162	
N1—H2N1···S4 ⁱ		0.91	2.47	3.359 (2)	165	
N1—H3N1···S1 ⁱⁱ		0.91	2.54	3.4297 (19)	165	
N1—H3N1···S3 ⁱⁱ		0.91	2.85	3.3019 (19)	112	
N2—H1N2···S3 ⁱⁱⁱ		0.91	2.58	3.376 (2)	147	
N2—H1N2···S2 ⁱⁱⁱ		0.91	2.92	3.580 (2)	131	
N2—H2N2…S4		0.91	2.43	3.309 (2)	164	
N2—H3N2···S1 ⁱ		0.91	2.50	3.393 (2)	169	
C2—H2A···S4 ⁱ		0.98	2.97	3.759 (3)	139	
C5—H5B···S3 ⁱⁱ		0.98	2.98	3.639 (6)	126	

Symmetry codes: (i) x, -y+1, z-1/2; (ii) -x+1/2, -y+3/2, -z+1; (iii) -x+1/2, y-1/2, -z+3/2.

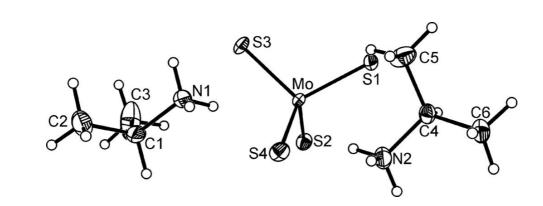
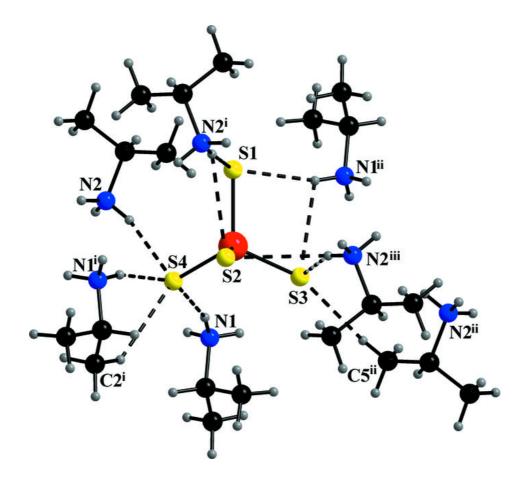


Fig. 1

Fig. 2



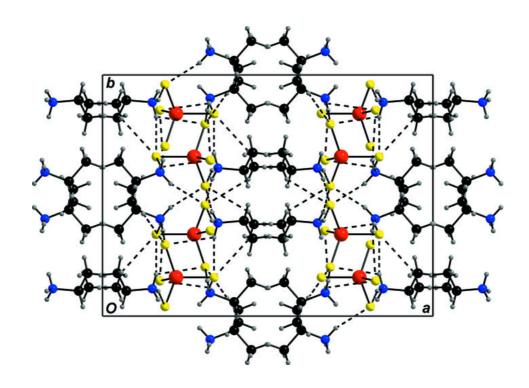


Fig. 3