

Acta Crystallographica Section E

Structure Reports

Online

ISSN 1600-5368

[2-(1-{2-[Azanidyl(ethylsulfanyl)methylidene- κ N]}hydrazin-1-ylidene- κ N¹}ethyl)-phenolato- κ O](dimethyl sulfoxide- κ O)-dioxidomolybdenum(VI)

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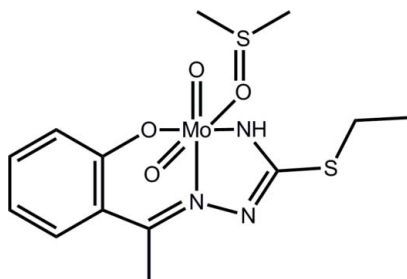
Received 11 June 2012; accepted 12 June 2012

Key indicators: single-crystal X-ray study; $T = 100$ K; mean $\sigma(\text{S}-\text{C}) = 0.003$ Å; disorder in main residue; R factor = 0.030; wR factor = 0.072; data-to-parameter ratio = 17.7.

The Mo^{VI} atom in the title complex, [Mo(C₁₁H₁₃N₃OS)₂O₂(C₂H₆OS)], is *N,N',O*-coordinated by the dianionic tridentate ligand, two mutually *cis* oxide O atoms and a dimethyl sulfoxide O atom, defining a distorted octahedral N₂O₄ donor set. The most prominent feature of the crystal packing is the formation of inversion dimers *via* pairs of N—H...O hydrogen bonds and eight-membered {...HNMoO}₂ loops. The Schiff base ligand is disordered over two orientations of equal occupancy.

Related literature

For the coordination chemistry and medicinal applications of thiosemicarbazone derivatives, see: Ahmadi *et al.* (2012); Dilworth & Hueting (2012). For related structures, see: Ceylan *et al.* (2009); Takjoo *et al.* (2012).



Experimental

Crystal data

[Mo(C₁₁H₁₃N₃OS)₂O₂(C₂H₆OS)]
 $M_r = 441.37$
 Triclinic, $P\bar{1}$
 $a = 8.0411$ (3) Å
 $b = 9.7243$ (3) Å
 $c = 12.0849$ (4) Å
 $\alpha = 73.387$ (3)°
 $\beta = 84.465$ (3)°
 $\gamma = 86.077$ (3)°
 $V = 900.47$ (5) Å³
 $Z = 2$
 Mo $K\alpha$ radiation
 $\mu = 0.98$ mm⁻¹
 $T = 100$ K
 $0.35 \times 0.15 \times 0.05$ mm

Data collection

Agilent SuperNova Dual diffractometer with an Atlas detector
 Absorption correction: multi-scan (*CrysAlis PRO*; Agilent, 2012)
 $T_{\min} = 0.691$, $T_{\max} = 1.000$
 13626 measured reflections
 4149 independent reflections
 3811 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.037$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.030$
 $wR(F^2) = 0.072$
 $S = 1.03$
 4149 reflections
 234 parameters
 100 restraints
 H-atom parameters constrained
 $\Delta\rho_{\max} = 0.73$ e Å⁻³
 $\Delta\rho_{\min} = -0.87$ e Å⁻³

Table 1

Selected bond lengths (Å).

Mo—O1	2.011 (7)	Mo—O4	1.7021 (18)
Mo—O2	2.2747 (16)	Mo—N1	2.193 (6)
Mo—O3	1.7133 (16)	Mo—N3	1.933 (15)

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>
N3—H3 <i>n</i> ...O3 ⁱ	0.88	2.23	3.090 (15)	166
N3'—H3 <i>n</i> '...O3 ⁱ	0.88	1.94	2.816 (16)	171

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

Data collection: *CrysAlis PRO* (Agilent, 2012); cell refinement: *CrysAlis PRO*; data reduction: *CrysAlis PRO*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3* (Farrugia, 1997) and *DIAMOND* (Brandenburg, 2006); software used to prepare material for publication: *publCIF* (Westrip, 2010).

The authors are grateful to the Ferdowsi University of Mashhad for financial support, and thank the Ministry of Higher Education (Malaysia) for funding structural studies through the High-Impact Research scheme (UM.C/HIR/MOHE/SC/3).

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB6848).

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supplementary materials

Acta Cryst. (2012). E68, m944–m945 [doi:10.1107/S1600536812026566]

[2-(1-{2-[Azanidyl(ethylsulfanyl)methylidene- κ N]}hydrazin-1-ylidene- κ N¹}ethyl)-phenolato- κ O](dimethyl sulfoxide- κ O)dioxidomolybdenum(VI)

Reza Takjoo, Seik Weng Ng and Edward R. T. Tiekink

Comment

Schiff bases derived from *S*-alkyl esters of thiosemicarbazone are capable of complexing both transition and main group metals (Ahmadi *et al.*, 2012) and these may be used as therapeutic and imaging agents (Dilworth & Hueting, 2012). Herein, the crystal and molecular structure of the title complex, (I), is described, which was determined as a part of on-going studies (Takjoo *et al.*, 2012).

The Mo^{VI} atom in (I), Fig. 1, exists within a distorted octahedral N₂O₄ donor set defined by the *N,N,O* atoms of the dinegative tridentate ligand, two oxo-O atoms and a DMSO-O atom, Table 1; the oxo-O atoms are *cis*. The dihedral angle between the five- and six-membered chelate rings is 11.34 (19) Å indicating some bending in the Schiff base ligand (the comparable angle in the second conformation of the disordered ligand is 7.81 (17)°). The molecular structure resembles that of the complex where the Mo atom is coordinated by methanol rather than DMSO (Ceylan *et al.*, 2009).

While whole molecule disorder of the Schiff base ligands precludes a detailed analysis of the crystal packing, a common feature of both orientations is the formation of N—H···O hydrogen bonds between centrosymmetrically related molecules to form a dimeric aggregate *via* an eight-membered {···HNMoO}₂ synthon, Fig. 2 and Table 2.

Experimental

An ethanolic solution (3 ml) of molybdenyl acetylacetonate (0.33 g, 1 mmol) was added drop-wise to an ethanolic solution (3 ml) of 1-(2-hydroxyphenyl)ethanone *S*-ethylisothiosemicarbazone hydrobromide (0.32 g, 1 mmol) under stirring. The clear solution was stirred for 1 h and yellow precipitate was appeared. The product was then filtered, washed with cold ethanol and dried in air. The resulting compound was dissolved in DMSO (2 ml) and by slow evaporation of the solvent orange prisms appeared after one week. *M.pt.* 427 K. Yield: 33%.

Refinement

Nitrogen- and carbon-bound H-atoms were placed in calculated positions [N—H = 0.88 Å and C—H = 0.95–0.99 Å, $U_{\text{iso}}(\text{H}) = 1.2\text{--}1.5U_{\text{eq}}(\text{N}, \text{C})$] and were included in the refinement in the riding model approximation.

The dianion is disordered over two positions in a 1:1 ratio. The benzene rings were refined as rigid hexagons of 1.39 Å sides and other pairs of bond distances were restrained to within 0.01 Å of each other. The anisotropic displacement parameters (restrained to be nearly isotropic) of the primed atoms were set to those of the unprimed ones.

Computing details

Data collection: *CrysAlis PRO* (Agilent, 2012); cell refinement: *CrysAlis PRO* (Agilent, 2012); data reduction: *CrysAlis PRO* (Agilent, 2012); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3* (Farrugia, 1997) and *DIAMOND* (Brandenburg,

2006); software used to prepare material for publication: *publCIF* (Westrip, 2010).

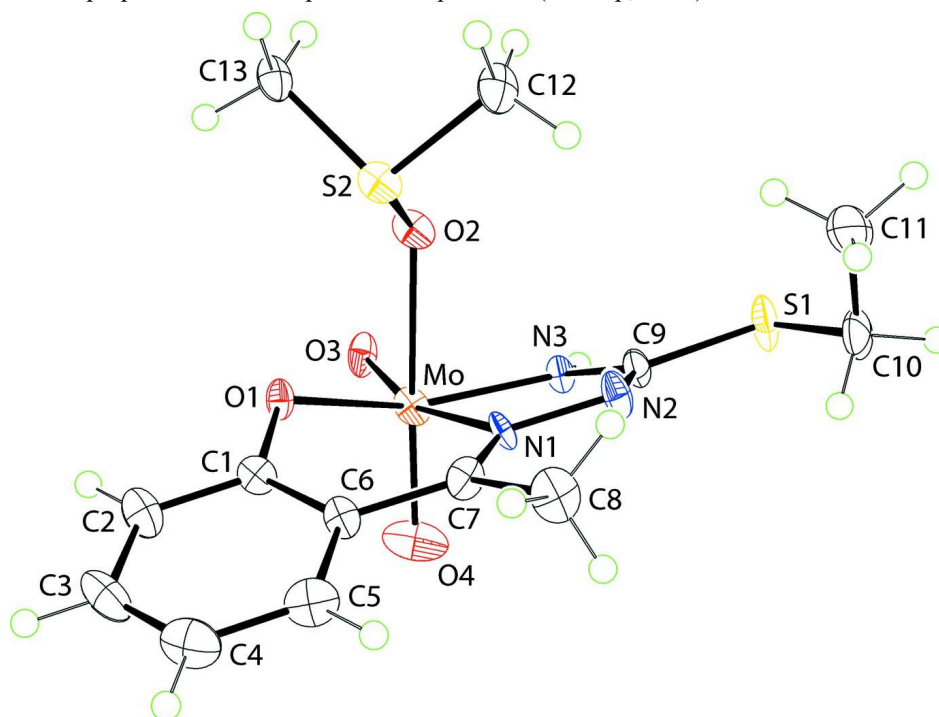


Figure 1

The molecular structure of (I) showing the atom-labelling scheme and displacement ellipsoids at the 50% probability level.

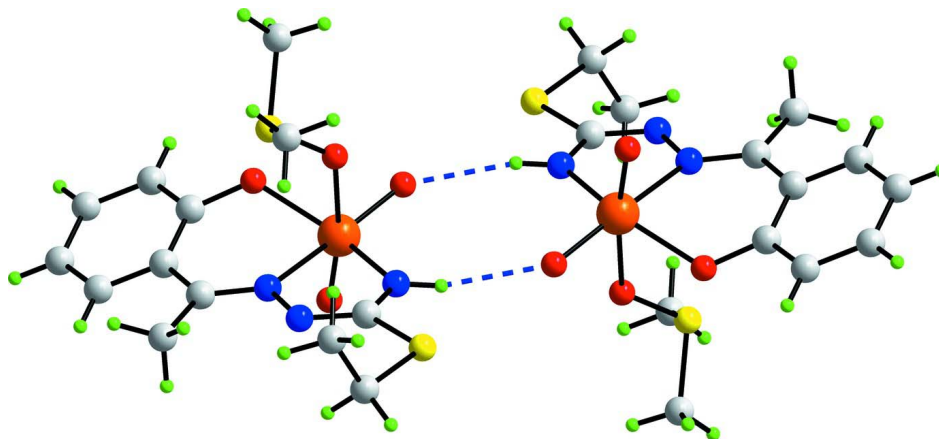


Figure 2

A view of the centrosymmetric aggregate in (I) mediated by N—H...O hydrogen bonds, shown as blue dashed lines. A similar arrangement is found for the second conformation of the Schiff base ligand.

[2-(1-{2-[Azanidyl(ethylsulfanyl)methylidene- κ N]}hydrazin-1-ylidene- κ N¹ethyl)phenolato- κ O](dimethyl sulfoxide- κ O)dioxidomolybdenum(VI)

Crystal data

[Mo(C ₁₁ H ₁₃ N ₃ OS)O ₂ (C ₂ H ₆ OS)]	Z = 2
$M_r = 441.37$	$F(000) = 448$
Triclinic, $P\bar{1}$	$D_x = 1.628 \text{ Mg m}^{-3}$
Hall symbol: -P 1	Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$
$a = 8.0411 (3) \text{ \AA}$	Cell parameters from 8498 reflections
$b = 9.7243 (3) \text{ \AA}$	$\theta = 2.4\text{--}27.5^\circ$
$c = 12.0849 (4) \text{ \AA}$	$\mu = 0.98 \text{ mm}^{-1}$
$\alpha = 73.387 (3)^\circ$	$T = 100 \text{ K}$
$\beta = 84.465 (3)^\circ$	Prism, orange
$\gamma = 86.077 (3)^\circ$	$0.35 \times 0.15 \times 0.05 \text{ mm}$
$V = 900.47 (5) \text{ \AA}^3$	

Data collection

Agilent SuperNova Dual	$T_{\min} = 0.691, T_{\max} = 1.000$
diffractometer with an Atlas detector	13626 measured reflections
Radiation source: SuperNova (Mo) X-ray	4149 independent reflections
Source	3811 reflections with $I > 2\sigma(I)$
Mirror monochromator	$R_{\text{int}} = 0.037$
Detector resolution: 10.4041 pixels mm^{-1}	$\theta_{\max} = 27.6^\circ, \theta_{\min} = 2.4^\circ$
ω scan	$h = -10 \rightarrow 10$
Absorption correction: multi-scan	$k = -12 \rightarrow 12$
(CrysAlis PRO; Agilent, 2012)	$l = -15 \rightarrow 15$

Refinement

Refinement on F^2	Secondary atom site location: difference Fourier
Least-squares matrix: full	map
$R[F^2 > 2\sigma(F^2)] = 0.030$	Hydrogen site location: inferred from
$wR(F^2) = 0.072$	neighbouring sites
$S = 1.03$	H-atom parameters constrained
4149 reflections	$w = 1/[\sigma^2(F_o^2) + (0.0278P)^2 + 1.0887P]$
234 parameters	where $P = (F_o^2 + 2F_c^2)/3$
100 restraints	$(\Delta/\sigma)_{\max} = 0.001$
Primary atom site location: structure-invariant	$\Delta\rho_{\max} = 0.73 \text{ e \AA}^{-3}$
direct methods	$\Delta\rho_{\min} = -0.87 \text{ e \AA}^{-3}$

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	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
Mo	0.39828 (2)	0.62362 (2)	0.284368 (16)	0.01779 (7)	
S2	-0.02453 (7)	0.59823 (6)	0.30257 (5)	0.02079 (13)	

O2	0.13555 (19)	0.56941 (17)	0.36645 (13)	0.0208 (3)	
O3	0.4236 (2)	0.67549 (16)	0.40557 (15)	0.0217 (3)	
O4	0.5828 (2)	0.6597 (2)	0.20227 (15)	0.0370 (5)	
C12	-0.1234 (3)	0.4305 (3)	0.3508 (2)	0.0268 (5)	
H12A	-0.0630	0.3620	0.3141	0.040*	
H12B	-0.2393	0.4441	0.3296	0.040*	
H12C	-0.1223	0.3934	0.4351	0.040*	
C13	-0.1570 (3)	0.6982 (2)	0.3810 (2)	0.0221 (5)	
H13A	-0.1176	0.7959	0.3631	0.033*	
H13B	-0.1547	0.6515	0.4643	0.033*	
H13C	-0.2717	0.7022	0.3588	0.033*	
S1	0.4023 (2)	0.14457 (15)	0.38075 (13)	0.0307 (3)	0.50
O1	0.2528 (12)	0.7850 (6)	0.1938 (4)	0.0162 (9)	0.50
N1	0.3150 (11)	0.5172 (5)	0.1633 (5)	0.0149 (10)	0.50
N2	0.3133 (9)	0.3667 (6)	0.2053 (4)	0.0203 (9)	0.50
N3	0.439 (3)	0.4201 (15)	0.3527 (10)	0.0137 (16)	0.50
H3n	0.4952	0.3893	0.4149	0.016*	0.50
C1	0.2566 (5)	0.8284 (3)	0.07631 (19)	0.0176 (9)	0.50
C2	0.2480 (5)	0.9757 (3)	0.0241 (3)	0.0275 (8)	0.50
H2	0.2412	1.0405	0.0703	0.033*	0.50
C3	0.2494 (5)	1.0281 (3)	-0.0957 (3)	0.0329 (9)	0.50
H3	0.2435	1.1287	-0.1314	0.040*	0.50
C4	0.2593 (5)	0.9332 (4)	-0.16331 (19)	0.0335 (9)	0.50
H4	0.2602	0.9691	-0.2452	0.040*	0.50
C5	0.2679 (5)	0.7860 (3)	-0.1111 (3)	0.0278 (8)	0.50
H5	0.2747	0.7211	-0.1573	0.033*	0.50
C6	0.2666 (4)	0.7335 (2)	0.0087 (3)	0.0183 (8)	0.50
C7	0.2706 (7)	0.5771 (6)	0.0576 (4)	0.0188 (9)	0.50
C8	0.2263 (7)	0.4788 (6)	-0.0121 (4)	0.0277 (8)	0.50
H8A	0.1657	0.3976	0.0391	0.042*	0.50
H8B	0.3291	0.4431	-0.0472	0.042*	0.50
H8C	0.1557	0.5326	-0.0732	0.042*	0.50
C9	0.3817 (12)	0.3257 (6)	0.3040 (5)	0.0185 (11)	0.50
C10	0.3280 (7)	0.0531 (6)	0.2844 (5)	0.0306 (9)	0.50
H10A	0.3694	0.1019	0.2038	0.037*	0.50
H10B	0.3759	-0.0467	0.3039	0.037*	0.50
C11	0.1370 (8)	0.0490 (7)	0.2911 (6)	0.0422 (12)	0.50
H11A	0.1011	0.0621	0.2134	0.063*	0.50
H11B	0.0866	0.1263	0.3222	0.063*	0.50
H11C	0.1013	-0.0439	0.3419	0.063*	0.50
S1'	0.3906 (2)	0.12666 (15)	0.43708 (13)	0.0307 (3)	0.50
O1'	0.2699 (12)	0.7538 (6)	0.1728 (4)	0.0162 (9)	0.50
N1'	0.3129 (11)	0.4692 (6)	0.1856 (5)	0.0149 (10)	0.50
N2'	0.3038 (9)	0.3236 (6)	0.2459 (4)	0.0203 (9)	0.50
N3'	0.434 (3)	0.4047 (15)	0.3800 (10)	0.0137 (16)	0.50
H3n'	0.4895	0.3818	0.4426	0.016*	0.50
C1'	0.2776 (5)	0.7739 (3)	0.05697 (19)	0.0176 (9)	0.50
C2'	0.2822 (5)	0.9143 (3)	-0.0140 (3)	0.0275 (8)	0.50
H2'	0.2816	0.9919	0.0193	0.033*	0.50

C3'	0.2878 (5)	0.9413 (3)	-0.1336 (3)	0.0329 (9)	0.50
H3'	0.2910	1.0372	-0.1821	0.040*	0.50
C4'	0.2888 (5)	0.8278 (4)	-0.18223 (19)	0.0335 (9)	0.50
H4'	0.2926	0.8462	-0.2640	0.040*	0.50
C5'	0.2841 (5)	0.6874 (3)	-0.1113 (3)	0.0278 (8)	0.50
H5'	0.2847	0.6099	-0.1445	0.033*	0.50
C6'	0.2785 (5)	0.6605 (3)	0.0083 (2)	0.0183 (8)	0.50
C7'	0.2681 (7)	0.5090 (6)	0.0789 (5)	0.0188 (9)	0.50
C8'	0.2100 (7)	0.3937 (6)	0.0307 (4)	0.0277 (8)	0.50
H8'A	0.1395	0.3280	0.0906	0.042*	0.50
H8'B	0.3076	0.3399	0.0065	0.042*	0.50
H8'C	0.1456	0.4388	-0.0362	0.042*	0.50
C9'	0.3720 (12)	0.3009 (6)	0.3435 (5)	0.0185 (11)	0.50
C10'	0.3169 (7)	0.0189 (5)	0.3535 (5)	0.0306 (9)	0.50
H10C	0.3611	0.0564	0.2717	0.037*	0.50
H10D	0.3621	-0.0809	0.3826	0.037*	0.50
C11'	0.1275 (8)	0.0176 (7)	0.3585 (6)	0.0422 (12)	0.50
H11D	0.0958	-0.0464	0.3151	0.063*	0.50
H11E	0.0821	0.1151	0.3241	0.063*	0.50
H11F	0.0823	-0.0167	0.4394	0.063*	0.50

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Mo	0.01666 (11)	0.01953 (11)	0.01794 (11)	-0.00370 (7)	-0.00961 (7)	-0.00315 (8)
S2	0.0173 (3)	0.0254 (3)	0.0192 (3)	-0.0038 (2)	-0.0098 (2)	-0.0022 (2)
O2	0.0160 (8)	0.0250 (8)	0.0199 (8)	-0.0045 (6)	-0.0099 (6)	-0.0005 (7)
O3	0.0200 (8)	0.0159 (8)	0.0345 (9)	0.0010 (6)	-0.0119 (7)	-0.0130 (7)
O4	0.0225 (9)	0.0648 (14)	0.0226 (9)	-0.0156 (9)	-0.0076 (7)	-0.0053 (9)
C12	0.0229 (12)	0.0233 (12)	0.0397 (14)	-0.0036 (10)	-0.0110 (10)	-0.0141 (11)
C13	0.0203 (11)	0.0164 (11)	0.0312 (12)	-0.0030 (9)	-0.0110 (9)	-0.0056 (9)
S1	0.0580 (6)	0.0097 (4)	0.0242 (6)	0.0007 (4)	-0.0099 (8)	-0.0029 (6)
O1	0.0220 (19)	0.012 (2)	0.0160 (17)	0.0021 (19)	-0.0077 (17)	-0.0041 (12)
N1	0.0204 (11)	0.008 (3)	0.015 (2)	-0.006 (3)	-0.0055 (18)	0.000 (2)
N2	0.0332 (15)	0.012 (3)	0.017 (3)	-0.005 (2)	-0.010 (2)	-0.0030 (19)
N3	0.0215 (15)	0.012 (2)	0.008 (5)	0.000 (2)	-0.009 (5)	-0.002 (3)
C1	0.0167 (15)	0.021 (2)	0.0157 (15)	-0.0021 (17)	-0.0070 (12)	-0.0037 (15)
C2	0.031 (2)	0.021 (2)	0.028 (2)	-0.0037 (16)	-0.0094 (16)	-0.0009 (13)
C3	0.036 (2)	0.028 (2)	0.028 (2)	-0.0096 (17)	-0.0079 (16)	0.0077 (15)
C4	0.029 (2)	0.045 (2)	0.0224 (18)	-0.0022 (18)	-0.0092 (15)	-0.0005 (18)
C5	0.0245 (17)	0.039 (2)	0.0215 (15)	-0.0002 (18)	-0.0089 (13)	-0.0087 (18)
C6	0.0172 (14)	0.017 (2)	0.0203 (14)	-0.0016 (19)	-0.0079 (11)	-0.003 (2)
C7	0.0164 (13)	0.021 (3)	0.025 (2)	-0.003 (2)	-0.0070 (14)	-0.014 (2)
C8	0.037 (2)	0.031 (2)	0.0202 (19)	-0.0053 (19)	-0.0133 (16)	-0.0112 (15)
C9	0.0272 (17)	0.011 (2)	0.018 (4)	-0.0034 (18)	-0.002 (3)	-0.004 (2)
C10	0.046 (2)	0.0160 (18)	0.034 (2)	-0.0063 (15)	0.000 (2)	-0.0140 (19)
C11	0.048 (2)	0.034 (3)	0.052 (3)	-0.0188 (19)	0.012 (3)	-0.026 (3)
S1'	0.0580 (6)	0.0097 (4)	0.0242 (6)	0.0007 (4)	-0.0099 (8)	-0.0029 (6)
O1'	0.0220 (19)	0.012 (2)	0.0160 (17)	0.0021 (19)	-0.0077 (17)	-0.0041 (12)

N1'	0.0204 (11)	0.008 (3)	0.015 (2)	-0.006 (3)	-0.0055 (18)	0.000 (2)
N2'	0.0332 (15)	0.012 (3)	0.017 (3)	-0.005 (2)	-0.010 (2)	-0.0030 (19)
N3'	0.0215 (15)	0.012 (2)	0.008 (5)	0.000 (2)	-0.009 (5)	-0.002 (3)
C1'	0.0167 (15)	0.021 (2)	0.0157 (15)	-0.0021 (17)	-0.0070 (12)	-0.0037 (15)
C2'	0.031 (2)	0.021 (2)	0.028 (2)	-0.0037 (16)	-0.0094 (16)	-0.0009 (13)
C3'	0.036 (2)	0.028 (2)	0.028 (2)	-0.0096 (17)	-0.0079 (16)	0.0077 (15)
C4'	0.029 (2)	0.045 (2)	0.0224 (18)	-0.0022 (18)	-0.0092 (15)	-0.0005 (18)
C5'	0.0245 (17)	0.039 (2)	0.0215 (15)	-0.0002 (18)	-0.0089 (13)	-0.0087 (18)
C6'	0.0172 (14)	0.017 (2)	0.0203 (14)	-0.0016 (19)	-0.0079 (11)	-0.003 (2)
C7'	0.0164 (13)	0.021 (3)	0.025 (2)	-0.003 (2)	-0.0070 (14)	-0.014 (2)
C8'	0.037 (2)	0.031 (2)	0.0202 (19)	-0.0053 (19)	-0.0133 (16)	-0.0112 (15)
C9'	0.0272 (17)	0.011 (2)	0.018 (4)	-0.0034 (18)	-0.002 (3)	-0.004 (2)
C10'	0.046 (2)	0.0160 (18)	0.034 (2)	-0.0063 (15)	0.000 (2)	-0.0140 (19)
C11'	0.048 (2)	0.034 (3)	0.052 (3)	-0.0188 (19)	0.012 (3)	-0.026 (3)

Geometric parameters (Å, °)

Mo—O1	2.011 (7)	C8—H8A	0.9800
Mo—O2	2.2747 (16)	C8—H8B	0.9800
Mo—O3	1.7133 (16)	C8—H8C	0.9800
Mo—O4	1.7021 (18)	C10—C11	1.533 (7)
Mo—O1'	1.897 (7)	C10—H10A	0.9900
Mo—N1	2.193 (6)	C10—H10B	0.9900
Mo—N3	1.933 (15)	C11—H11A	0.9800
Mo—N3'	2.127 (15)	C11—H11B	0.9800
Mo—N1'	2.340 (6)	C11—H11C	0.9800
S2—O2	1.5330 (15)	S1'—C9'	1.753 (5)
S2—C13	1.780 (2)	S1'—C10'	1.810 (5)
S2—C12	1.782 (2)	O1'—C1'	1.353 (5)
C12—H12A	0.9800	N1'—C7'	1.315 (6)
C12—H12B	0.9800	N1'—N2'	1.398 (5)
C12—H12C	0.9800	N2'—C9'	1.304 (6)
C13—H13A	0.9800	N3'—C9'	1.353 (6)
C13—H13B	0.9800	N3'—H3n'	0.8800
C13—H13C	0.9800	C1'—C2'	1.3900
S1—C9	1.746 (6)	C1'—C6'	1.3900
S1—C10	1.818 (5)	C2'—C3'	1.3900
O1—C1	1.359 (5)	C2'—H2'	0.9500
N1—C7	1.316 (6)	C3'—C4'	1.3900
N1—N2	1.406 (5)	C3'—H3'	0.9500
N2—C9	1.307 (6)	C4'—C5'	1.3900
N3—C9	1.351 (6)	C4'—H4'	0.9500
N3—H3n	0.8800	C5'—C6'	1.3900
C1—C2	1.3900	C5'—H5'	0.9500
C1—C6	1.3900	C6'—C7'	1.479 (5)
C2—C3	1.3900	C7'—C8'	1.521 (6)
C2—H2	0.9500	C8'—H8'A	0.9800
C3—C4	1.3900	C8'—H8'B	0.9800
C3—H3	0.9500	C8'—H8'C	0.9800
C4—C5	1.3900	C10'—C11'	1.519 (7)

C4—H4	0.9500	C10'—H10C	0.9900
C5—C6	1.3900	C10'—H10D	0.9900
C5—H5	0.9500	C11'—H11D	0.9800
C6—C7	1.465 (6)	C11'—H11E	0.9800
C7—C8	1.522 (6)	C11'—H11F	0.9800
O4—Mo—O3	104.39 (8)	C6—C5—H5	120.0
O4—Mo—O1'	94.2 (3)	C4—C5—H5	120.0
O3—Mo—O1'	115.50 (14)	C5—C6—C1	120.0
O4—Mo—N3	98.5 (7)	C5—C6—C7	116.8 (3)
O3—Mo—N3	96.7 (3)	C1—C6—C7	123.2 (3)
O1'—Mo—N3	141.2 (5)	N1—C7—C6	120.7 (4)
O4—Mo—O1	99.5 (3)	N1—C7—C8	117.9 (5)
O3—Mo—O1	102.65 (14)	C6—C7—C8	121.4 (4)
O1'—Mo—O1	12.9 (2)	N2—C9—N3	122.3 (8)
N3—Mo—O1	149.3 (6)	N2—C9—S1	121.5 (5)
O4—Mo—N3'	103.0 (6)	N3—C9—S1	116.2 (8)
O3—Mo—N3'	90.1 (3)	C11—C10—S1	113.6 (4)
O1'—Mo—N3'	144.7 (5)	C11—C10—H10A	108.9
N3—Mo—N3'	7.2 (8)	S1—C10—H10A	108.9
O1—Mo—N3'	150.4 (6)	C11—C10—H10B	108.9
O4—Mo—N1	90.4 (2)	S1—C10—H10B	108.9
O3—Mo—N1	163.71 (19)	H10A—C10—H10B	107.7
O1'—Mo—N1	69.3 (2)	C10—C11—H11A	109.5
N3—Mo—N1	74.0 (3)	C10—C11—H11B	109.5
O1—Mo—N1	81.2 (2)	H11A—C11—H11B	109.5
N3'—Mo—N1	79.8 (3)	C10—C11—H11C	109.5
O4—Mo—O2	170.16 (7)	H11A—C11—H11C	109.5
O3—Mo—O2	85.28 (7)	H11B—C11—H11C	109.5
O1'—Mo—O2	79.7 (3)	C9'—S1'—C10'	102.2 (3)
N3—Mo—O2	82.0 (7)	C1'—O1'—Mo	128.9 (4)
O1—Mo—O2	76.2 (3)	C7'—N1'—N2'	117.3 (5)
N3'—Mo—O2	78.5 (6)	C7'—N1'—Mo	125.2 (4)
N1—Mo—O2	80.3 (2)	N2'—N1'—Mo	117.5 (4)
O4—Mo—N1'	94.7 (2)	C9'—N2'—N1'	108.8 (5)
O3—Mo—N1'	154.27 (16)	C9'—N3'—Mo	119.4 (9)
O1'—Mo—N1'	79.4 (2)	C9'—N3'—H3n'	120.3
N3—Mo—N1'	63.1 (3)	Mo—N3'—H3n'	120.3
O1—Mo—N1'	90.7 (2)	O1'—C1'—C2'	117.6 (3)
N3'—Mo—N1'	68.7 (3)	O1'—C1'—C6'	122.4 (3)
N1—Mo—N1'	11.27 (17)	C2'—C1'—C6'	120.0
O2—Mo—N1'	76.6 (2)	C1'—C2'—C3'	120.0
O2—S2—C13	103.33 (10)	C1'—C2'—H2'	120.0
O2—S2—C12	103.52 (10)	C3'—C2'—H2'	120.0
C13—S2—C12	99.67 (12)	C4'—C3'—C2'	120.0
S2—O2—Mo	125.84 (9)	C4'—C3'—H3'	120.0
S2—C12—H12A	109.5	C2'—C3'—H3'	120.0
S2—C12—H12B	109.5	C3'—C4'—C5'	120.0
H12A—C12—H12B	109.5	C3'—C4'—H4'	120.0

S2—C12—H12C	109.5	C5'—C4'—H4'	120.0
H12A—C12—H12C	109.5	C6'—C5'—C4'	120.0
H12B—C12—H12C	109.5	C6'—C5'—H5'	120.0
S2—C13—H13A	109.5	C4'—C5'—H5'	120.0
S2—C13—H13B	109.5	C5'—C6'—C1'	120.0
H13A—C13—H13B	109.5	C5'—C6'—C7'	117.2 (3)
S2—C13—H13C	109.5	C1'—C6'—C7'	122.7 (3)
H13A—C13—H13C	109.5	N1'—C7'—C6'	120.5 (4)
H13B—C13—H13C	109.5	N1'—C7'—C8'	117.6 (5)
C9—S1—C10	103.3 (3)	C6'—C7'—C8'	121.8 (4)
C1—O1—Mo	124.2 (4)	C7'—C8'—H8'A	109.5
C7—N1—N2	117.6 (5)	C7'—C8'—H8'B	109.5
C7—N1—Mo	127.8 (4)	H8'A—C8'—H8'B	109.5
N2—N1—Mo	114.5 (4)	C7'—C8'—H8'C	109.5
C9—N2—N1	108.9 (5)	H8'A—C8'—H8'C	109.5
C9—N3—Mo	119.3 (10)	H8'B—C8'—H8'C	109.5
C9—N3—H3n	120.3	N2'—C9'—N3'	124.3 (8)
Mo—N3—H3n	120.3	N2'—C9'—S1'	120.4 (5)
O1—C1—C2	116.7 (3)	N3'—C9'—S1'	115.2 (7)
O1—C1—C6	123.3 (3)	C11'—C10'—S1'	113.5 (4)
C2—C1—C6	120.0	C11'—C10'—H10C	108.9
C3—C2—C1	120.0	S1'—C10'—H10C	108.9
C3—C2—H2	120.0	C11'—C10'—H10D	108.9
C1—C2—H2	120.0	S1'—C10'—H10D	108.9
C4—C3—C2	120.0	H10C—C10'—H10D	107.7
C4—C3—H3	120.0	C10'—C11'—H11D	109.5
C2—C3—H3	120.0	C10'—C11'—H11E	109.5
C3—C4—C5	120.0	H11D—C11'—H11E	109.5
C3—C4—H4	120.0	C10'—C11'—H11F	109.5
C5—C4—H4	120.0	H11D—C11'—H11F	109.5
C6—C5—C4	120.0	H11E—C11'—H11F	109.5
C13—S2—O2—Mo	-125.78 (12)	Mo—N3—C9—S1	-172.6 (9)
C12—S2—O2—Mo	130.66 (13)	C10—S1—C9—N2	4.7 (8)
O3—Mo—O2—S2	139.32 (13)	C10—S1—C9—N3	-175.7 (12)
O1'—Mo—O2—S2	22.32 (18)	C9—S1—C10—C11	-81.4 (5)
N3—Mo—O2—S2	-123.3 (3)	O4—Mo—O1'—C1'	44.6 (7)
O1—Mo—O2—S2	35.04 (18)	O3—Mo—O1'—C1'	152.7 (6)
N3'—Mo—O2—S2	-129.7 (3)	N3—Mo—O1'—C1'	-64.4 (14)
N1—Mo—O2—S2	-48.22 (18)	O1—Mo—O1'—C1'	160 (3)
N1'—Mo—O2—S2	-59.11 (17)	N3'—Mo—O1'—C1'	-74.9 (13)
O4—Mo—O1—C1	42.4 (7)	N1—Mo—O1'—C1'	-44.2 (7)
O3—Mo—O1—C1	149.6 (6)	O2—Mo—O1'—C1'	-127.6 (8)
O1'—Mo—O1—C1	-23.9 (18)	N1'—Mo—O1'—C1'	-49.4 (7)
N3—Mo—O1—C1	-82.8 (13)	O4—Mo—N1'—C7'	-69.8 (7)
N3'—Mo—O1—C1	-97.0 (11)	O3—Mo—N1'—C7'	151.9 (4)
N1—Mo—O1—C1	-46.5 (7)	O1'—Mo—N1'—C7'	23.5 (7)
O2—Mo—O1—C1	-128.6 (7)	N3—Mo—N1'—C7'	-166.9 (11)
N1'—Mo—O1—C1	-52.6 (7)	O1—Mo—N1'—C7'	29.8 (8)

O4—Mo—N1—C7	-71.6 (8)	N3'—Mo—N1'—C7'	-171.9 (11)
O3—Mo—N1—C7	133.1 (5)	N1—Mo—N1'—C7'	-2.3 (17)
O1'—Mo—N1—C7	22.7 (7)	O2—Mo—N1'—C7'	105.4 (7)
N3—Mo—N1—C7	-170.3 (11)	O4—Mo—N1'—N2'	112.3 (6)
O1—Mo—N1—C7	28.0 (8)	O3—Mo—N1'—N2'	-26.0 (10)
N3'—Mo—N1—C7	-174.7 (10)	O1'—Mo—N1'—N2'	-154.4 (7)
O2—Mo—N1—C7	105.3 (8)	N3—Mo—N1'—N2'	15.2 (9)
N1'—Mo—N1—C7	175 (3)	O1—Mo—N1'—N2'	-148.1 (7)
O4—Mo—N1—N2	107.1 (6)	N3'—Mo—N1'—N2'	10.2 (9)
O3—Mo—N1—N2	-48.2 (12)	N1—Mo—N1'—N2'	180 (3)
O1'—Mo—N1—N2	-158.6 (7)	O2—Mo—N1'—N2'	-72.5 (6)
N3—Mo—N1—N2	8.4 (9)	C7'—N1'—N2'—C9'	172.1 (8)
O1—Mo—N1—N2	-153.3 (7)	Mo—N1'—N2'—C9'	-9.8 (9)
N3'—Mo—N1—N2	4.0 (9)	O4—Mo—N3'—C9'	-98.7 (15)
O2—Mo—N1—N2	-76.0 (6)	O3—Mo—N3'—C9'	156.5 (15)
N1'—Mo—N1—N2	-5.9 (18)	O1'—Mo—N3'—C9'	18 (2)
C7—N1—N2—C9	171.6 (8)	N3—Mo—N3'—C9'	-47 (8)
Mo—N1—N2—C9	-7.2 (9)	O1—Mo—N3'—C9'	40 (2)
O4—Mo—N3—C9	-95.9 (15)	N1—Mo—N3'—C9'	-10.7 (14)
O3—Mo—N3—C9	158.4 (14)	O2—Mo—N3'—C9'	71.4 (15)
O1'—Mo—N3—C9	12 (2)	N1'—Mo—N3'—C9'	-8.6 (13)
O1—Mo—N3—C9	30 (2)	Mo—O1'—C1'—C2'	-132.5 (5)
N3'—Mo—N3—C9	135 (10)	Mo—O1'—C1'—C6'	48.3 (9)
N1—Mo—N3—C9	-7.9 (13)	O1'—C1'—C2'—C3'	-179.1 (6)
O2—Mo—N3—C9	74.2 (15)	C6'—C1'—C2'—C3'	0.0
N1'—Mo—N3—C9	-4.8 (12)	C1'—C2'—C3'—C4'	0.0
Mo—O1—C1—C2	-138.2 (4)	C2'—C3'—C4'—C5'	0.0
Mo—O1—C1—C6	42.6 (9)	C3'—C4'—C5'—C6'	0.0
O1—C1—C2—C3	-179.3 (6)	C4'—C5'—C6'—C1'	0.0
C6—C1—C2—C3	0.0	C4'—C5'—C6'—C7'	178.0 (4)
C1—C2—C3—C4	0.0	O1'—C1'—C6'—C5'	179.1 (6)
C2—C3—C4—C5	0.0	C2'—C1'—C6'—C5'	0.0
C3—C4—C5—C6	0.0	O1'—C1'—C6'—C7'	1.2 (6)
C4—C5—C6—C1	0.0	C2'—C1'—C6'—C7'	-177.9 (4)
C4—C5—C6—C7	178.1 (4)	N2'—N1'—C7'—C6'	-178.4 (6)
O1—C1—C6—C5	179.2 (6)	Mo—N1'—C7'—C6'	3.7 (10)
C2—C1—C6—C5	0.0	N2'—N1'—C7'—C8'	0.1 (11)
O1—C1—C6—C7	1.3 (6)	Mo—N1'—C7'—C8'	-177.8 (5)
C2—C1—C6—C7	-178.0 (4)	C5'—C6'—C7'—N1'	159.1 (6)
N2—N1—C7—C6	179.0 (6)	C1'—C6'—C7'—N1'	-23.0 (8)
Mo—N1—C7—C6	-2.4 (11)	C5'—C6'—C7'—C8'	-19.4 (6)
N2—N1—C7—C8	-2.0 (11)	C1'—C6'—C7'—C8'	158.6 (4)
Mo—N1—C7—C8	176.7 (5)	N1'—N2'—C9'—N3'	2.3 (17)
C5—C6—C7—N1	161.3 (6)	N1'—N2'—C9'—S1'	-176.8 (6)
C1—C6—C7—N1	-20.6 (8)	Mo—N3'—C9'—N2'	7 (2)
C5—C6—C7—C8	-17.7 (6)	Mo—N3'—C9'—S1'	-173.7 (8)
C1—C6—C7—C8	160.3 (4)	C10'—S1'—C9'—N2'	5.2 (8)
N1—N2—C9—N3	0.9 (16)	C10'—S1'—C9'—N3'	-174.0 (12)
N1—N2—C9—S1	-179.5 (6)	C9'—S1'—C10'—C11'	-80.7 (5)

Mo—N3—C9—N2 7 (2)

Hydrogen-bond geometry (Å, °)

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
N3—H3 <i>n</i> \cdots O3 ⁱ	0.88	2.23	3.090 (15)	166
N3'—H3 <i>n'</i> \cdots O3 ⁱ	0.88	1.94	2.816 (16)	171

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