

(4,4',6,6'-Tetra-*tert*-butyl-2,2'-{[2-(di-methylamino)ethyl]nitrilobis(methylene)}diphenolato)dioxidomolybdenum(VI) chloroform monosolvate

Xiangyang Lei* and Nagasree Chelamalla

Department of Chemistry & Biochemistry, Lamar University, Beaumont, TX 77710,

USA

Correspondence e-mail: xlei@lamar.edu

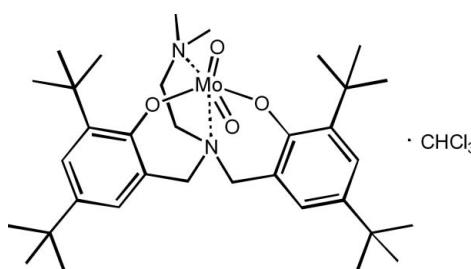
Received 12 August 2011; accepted 4 October 2011

Key indicators: single-crystal X-ray study; $T = 110\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.006\text{ \AA}$; R factor = 0.031; wR factor = 0.078; data-to-parameter ratio = 13.2.

In the title compound, $[\text{Mo}(\text{C}_{34}\text{H}_{54}\text{N}_2\text{O}_2)\text{O}_2]\cdot\text{CHCl}_3$, the molybdenum(VI) ion exhibits a *cis*-dioxide distorted octahedral geometry. Two anionic phenolate O-atom donors and two neutral N-atom donors of the ligand are *trans* and *cis*, respectively. The Mo=O bond lengths and the O=Mo=O bond angle are typical for six-coordinated dioxomolybdenum(VI) complexes. The Mo–N bond lengths are longer than 2.30 Å, as expected for a *trans* O=Mo=N structure.

Related literature

For molybdenum coordination complexes as catalysts, see: Wong *et al.* (2010); Rappe & Goddard (1982). For the synthesis of the ligand, see: Tshuva *et al.* (2001). For incorporation of the molybdenum center into redox enzymes, see: Tucci *et al.* (1998); Schultz *et al.* (1993). For spectroscopic and NMR data, see: Lehtonen *et al.* (2006). For related structures, see: Hinshaw *et al.* (1989); Lehtonen & Sillanpää (2005).



Experimental

Crystal data

$[\text{Mo}(\text{C}_{34}\text{H}_{54}\text{N}_2\text{O}_2)\text{O}_2]\cdot\text{CHCl}_3$

$M_r = 770.10$

Orthorhombic, $Pna2_1$
 $a = 24.3475(10)\text{ \AA}$
 $b = 13.9748(6)\text{ \AA}$
 $c = 11.0267(4)\text{ \AA}$
 $V = 3751.9(3)\text{ \AA}^3$

$Z = 4$
Cu $K\alpha$ radiation
 $\mu = 5.12\text{ mm}^{-1}$
 $T = 110\text{ K}$
 $0.50 \times 0.20 \times 0.02\text{ mm}$

Data collection

Bruker MWPC area-detector diffractometer
Absorption correction: multi-scan (*SADABS*; Bruker, 2008)
 $R_{\text{int}} = 0.081$
 $T_{\text{min}} = 0.184$, $T_{\text{max}} = 0.904$

79926 measured reflections
5548 independent reflections
5061 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.081$
 $\theta_{\text{max}} = 60.0^\circ$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.031$
 $wR(F^2) = 0.078$
 $S = 1.00$
5548 reflections
421 parameters
1 restraint

H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.81\text{ e \AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.51\text{ e \AA}^{-3}$
Absolute structure: Flack (1983),
2649 Friedel pairs
Flack parameter: 0.000 (9)

Data collection: *FRAMBO* (Bruker, 1999); cell refinement: *FRAMBO*; data reduction: *SAINT* (Bruker, 2004); 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*.

We thank the Welch Foundation (V-004) for financial support. We are very grateful to Dr Joseph Reibenspies at Texas A & M University for the X-ray crystallographic analysis. The X-ray diffractometers, small angle scattering instrumentation and crystallographic computing systems in the X-ray Diffraction Laboratory at the Department of Chemistry, Texas A & M University were purchased with funds provided by the National Science Foundation (CHE-9807975, CHE-0079822 and CHE-0215838).

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: JJ2100).

References

- Bruker (1999). *FRAMBO*. Bruker AXS Inc., Madison, Wisconsin, USA.
- Bruker (2004). *SAINT*. Bruker AXS Inc., Madison, Wisconsin, USA.
- Bruker (2008). *SADABS*. Bruker AXS Inc., Madison, Wisconsin, USA.
- Flack, H. D. (1983). *Acta Cryst. A39*, 876–881.
- Hinshaw, C. J., Peng, G., Singh, R., Spence, J. T., Enemark, J. H., Bruck, M., Kristofzski, J., Merbs, S. L., Ortega, R. B. & Wexler, P. A. (1989). *Inorg. Chem.* **28**, 4483–4491.
- Lehtonen, A. & Sillanpää, R. (2005). *Polyhedron*, **24**, 257–265.
- Lehtonen, A., Wasberg, M. & Sillanpää, R. (2006). *Polyhedron*, **25**, 767–775.
- Rappe, A. K. & Goddard, W. A. (1982). *J. Am. Chem. Soc.* **104**, 448–456.
- Schultz, B. E., Gheller, S. F., Muetterties, M. C., Scott, M. J. & Holm, R. H. (1993). *J. Am. Chem. Soc.* **115**, 2714–2722.
- Sheldrick, G. M. (2008). *Acta Cryst. A64*, 112–122.
- Tshuva, E., Goldberg, I. & Kol, M. (2001). *Organometallics*, **20**, 3017–3028.
- Tucci, G. C., Donahue, J. P. & Holm, R. H. (1998). *Inorg. Chem.* **37**, 1602–1608.
- Wong, Y. L., Tong, L. H., Dilworth, J. R., Ng, D. K. P. & Lee, H. K. (2010). *Dalton Trans.* **39**, 4602–4611.

supporting information

Acta Cryst. (2011). E67, m1510 [doi:10.1107/S160053681104092X]

(4,4',6,6'-Tetra-*tert*-butyl-2,2'-{[2-(dimethylamino)ethyl]nitrilobis(methylene)}diphenolato)dioxidomolybdenum(VI) chloroform monosolvate

Xiangyang Lei and Nagasree Chelamalla

S1. Comment

Molybdenum coordination complexes have attracted considerable attention because they can catalyze a variety of chemical reactions such as olefin epoxidation (Wong *et al.* 2010) and olefin metathesis (Rappe & Goddard, 1982) reactions. In addition, molybdenum is also a necessary element in diverse biological systems whereby the molybdenum center is incorporated into various redox enzymes such as DMSO reductase (Tucci *et al.* 1998) and xanthine oxidase (Schultz *et al.* 1993). A number of related dioxomolybdenum(VI) complexes with tetradentate ligands have been reported (Hinshaw *et al.* 1989; Lehtonen & Sillanpää, 2005).

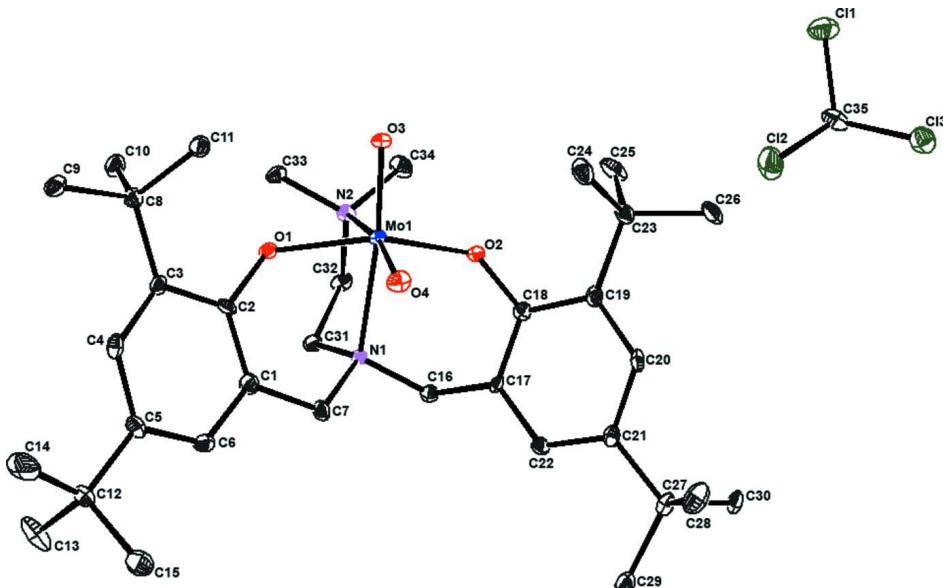
While the X-ray structure of the title compound is described here, its synthesis, IR, and ^1H & ^{13}C NMR data have been reported (Lehtonen *et al.* 2006). The title complex contains one crystallographically unique molybdenum ion in a *cis*-dioxo distorted octahedral geometry. The aminobis(phenolate) moiety is coordinated to the MoO_2^{2+} unit as a tripodal tetradentate ligand through two anionic phenolate oxygen donors (*trans* to each other) and two neutral nitrogen donors (*cis* to each other). The Mo=O bond lengths (1.702 (2) and 1.702 (3) Å for Mo=O3 and Mo=O4, respectively) and the O=Mo=O bond angle (108.33 (13) $^\circ$) are typical for six-coordinated dioxomolybdenum(VI) complexes. The bond lengths of Mo—N1 and Mo—N2 are 2.392 (3) and 2.422 (3) Å, respectively, both of which are > 2.30 Å as expected for the *trans* O=Mo—N structure as well as a distorted octahedral geometry.

S2. Experimental

To a solution of 0.52 g (1.00 mmol) of 6,6'-(2-(dimethylamino)ethylazanediyi)bis(methylene)bis(2,4-di-*tert*-butylphenol) (Tshuva *et al.* 2001) in 10 ml of CH_2Cl_2 and 10 ml of CH_3OH was added 0.33 g (1.06 mmol) of $\text{MoO}_2(\text{acac})_2$. The resulting orange solution was stirred overnight at room temperature. The yellow solid (0.61 g) was collected by filtration and washed with cold methanol. Single crystals suitable for X-ray diffraction were obtained by recrystallization from $\text{CHCl}_3/\text{hexanes}$.

S3. Refinement

All non-hydrogen atoms were refined with anisotropic thermal parameters. The hydrogen atoms bound to carbon atoms were placed in idealized positions and constrained to ride on their parent atoms, with $d(\text{C}—\text{H}) = 0.95\text{--}1.00$ Å, $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$. The number of Friedel pairs used for absolute structure refinement is 2649.

**Figure 1**

A view of the molecular structure. Ellipsoids are drawn at the 30% probability level. Hydrogen atoms were omitted for clarity.

(4,4',6,6'-Tetra-tert-butyl-2,2'-(dimethylamino)ethyl)nitrilobis(methylene)diphenolato)dioxidomolybdenum(VI) chloroform monosolvate

Crystal data



$M_r = 770.10$

Orthorhombic, $Pna2_1$

Hall symbol: P 2c -2n

$a = 24.3475 (10)$ Å

$b = 13.9748 (6)$ Å

$c = 11.0267 (4)$ Å

$V = 3751.9 (3)$ Å³

$Z = 4$

$F(000) = 1616$

$D_x = 1.363 \text{ Mg m}^{-3}$

Cu $K\alpha$ radiation, $\lambda = 1.54178$ Å

Cell parameters from 9879 reflections

$\theta = 3.6\text{--}62.7^\circ$

$\mu = 5.12 \text{ mm}^{-1}$

$T = 110$ K

Plate, yellow

$0.50 \times 0.20 \times 0.02$ mm

Data collection

Bruker MWPC area-detector
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

φ and ω scans

Absorption correction: multi-scan
(*SADABS*; Bruker, 2008)

$T_{\min} = 0.184$, $T_{\max} = 0.904$

79926 measured reflections

5548 independent reflections

5061 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.081$

$\theta_{\max} = 60.0^\circ$, $\theta_{\min} = 4.8^\circ$

$h = -27 \rightarrow 27$

$k = -15 \rightarrow 15$

$l = -12 \rightarrow 12$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.031$

$wR(F^2) = 0.078$

$S = 1.00$

5548 reflections

421 parameters

1 restraint

Primary atom site location: structure-invariant
direct methods

Secondary atom site location: difference Fourier map

Hydrogen site location: inferred from neighbouring sites

H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.054P)^2]$$

$$\text{where } P = (F_o^2 + 2F_c^2)/3$$

$$(\Delta/\sigma)_{\max} = 0.001$$

$$\Delta\rho_{\max} = 0.81 \text{ e \AA}^{-3}$$

$$\Delta\rho_{\min} = -0.51 \text{ e \AA}^{-3}$$

Absolute structure: Flack (1983), 2649 Friedel pairs

Absolute structure parameter: 0.000 (9)

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 $F^2 > 2\sigma(F^2)$ 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	$U_{\text{iso}}^* / U_{\text{eq}}$
Mo1	0.426810 (10)	0.838846 (17)	-0.50398 (3)	0.01403 (9)
C11	0.59086 (5)	0.19093 (10)	-0.37928 (12)	0.0456 (3)
Cl2	0.62313 (6)	0.38887 (9)	-0.37779 (18)	0.0700 (5)
Cl3	0.70222 (5)	0.24265 (8)	-0.43262 (11)	0.0380 (3)
O1	0.37192 (9)	0.93850 (17)	-0.4855 (3)	0.0161 (6)
O2	0.48994 (10)	0.76272 (18)	-0.4523 (3)	0.0198 (6)
O3	0.38528 (10)	0.74922 (18)	-0.5542 (2)	0.0195 (6)
O4	0.45879 (11)	0.8885 (2)	-0.6260 (2)	0.0223 (6)
N1	0.47151 (12)	0.9508 (2)	-0.3729 (3)	0.0135 (7)
N2	0.39503 (13)	0.8003 (2)	-0.3016 (3)	0.0171 (7)
C1	0.42256 (14)	1.0851 (3)	-0.4926 (5)	0.0205 (9)
C2	0.37556 (14)	1.0319 (2)	-0.5192 (4)	0.0176 (9)
C3	0.33115 (15)	1.0736 (3)	-0.5830 (3)	0.0159 (9)
C4	0.33686 (16)	1.1685 (3)	-0.6165 (4)	0.0188 (9)
H4	0.3075	1.1973	-0.6600	0.023*
C5	0.38261 (16)	1.2249 (3)	-0.5909 (4)	0.0198 (9)
C6	0.42495 (15)	1.1812 (3)	-0.5263 (4)	0.0202 (11)
H6	0.4563	1.2179	-0.5047	0.024*
C7	0.47454 (16)	1.0452 (3)	-0.4337 (4)	0.0199 (9)
H7A	0.5032	1.0411	-0.4972	0.024*
H7B	0.4873	1.0924	-0.3731	0.024*
C8	0.27985 (16)	1.0155 (3)	-0.6168 (4)	0.0184 (9)
C9	0.23863 (16)	1.0749 (3)	-0.6897 (4)	0.0230 (9)
H9A	0.2064	1.0356	-0.7085	0.035*
H9B	0.2558	1.0964	-0.7653	0.035*
H9BC	0.2273	1.1306	-0.6419	0.035*
C10	0.24975 (14)	0.9810 (3)	-0.5024 (5)	0.0239 (8)
H10A	0.2181	0.9419	-0.5260	0.036*
H10B	0.2371	1.0364	-0.4557	0.036*

H10C	0.2748	0.9426	-0.4528	0.036*
C11	0.29541 (17)	0.9285 (3)	-0.6952 (4)	0.0213 (9)
H11A	0.2619	0.8972	-0.7245	0.032*
H11B	0.3168	0.8831	-0.6465	0.032*
H11C	0.3174	0.9498	-0.7645	0.032*
C12	0.38648 (18)	1.3300 (3)	-0.6294 (4)	0.0252 (10)
C13	0.3754 (2)	1.3935 (3)	-0.5203 (5)	0.0456 (14)
H13A	0.3805	1.4606	-0.5432	0.068*
H13B	0.4010	1.3771	-0.4549	0.068*
H13C	0.3376	1.3836	-0.4924	0.068*
C14	0.3457 (2)	1.3543 (3)	-0.7301 (5)	0.0437 (14)
H14A	0.3502	1.4215	-0.7537	0.066*
H14B	0.3082	1.3439	-0.7008	0.066*
H14C	0.3526	1.3131	-0.8004	0.066*
C15	0.4439 (2)	1.3524 (3)	-0.6790 (5)	0.0355 (12)
H15A	0.4457	1.4198	-0.7031	0.053*
H15B	0.4513	1.3118	-0.7496	0.053*
H15C	0.4714	1.3399	-0.6161	0.053*
C16	0.52969 (15)	0.9236 (3)	-0.3376 (4)	0.0192 (9)
H16A	0.5495	0.9822	-0.3125	0.023*
H16B	0.5281	0.8804	-0.2665	0.023*
C17	0.56204 (14)	0.8750 (3)	-0.4370 (4)	0.0170 (9)
C18	0.54237 (14)	0.7891 (3)	-0.4836 (4)	0.0178 (9)
C19	0.57446 (15)	0.7309 (3)	-0.5608 (4)	0.0190 (9)
C20	0.62607 (15)	0.7660 (3)	-0.5902 (4)	0.0176 (9)
H20	0.6489	0.7279	-0.6407	0.021*
C21	0.64651 (16)	0.8541 (3)	-0.5501 (4)	0.0174 (9)
C22	0.61418 (15)	0.9075 (3)	-0.4715 (3)	0.0182 (10)
H22	0.6275	0.9666	-0.4409	0.022*
C23	0.55331 (17)	0.6345 (3)	-0.6097 (4)	0.0217 (9)
C24	0.5097 (2)	0.6521 (3)	-0.7063 (5)	0.0348 (12)
H24A	0.4975	0.5908	-0.7399	0.052*
H24B	0.5252	0.6916	-0.7713	0.052*
H24C	0.4783	0.6853	-0.6699	0.052*
C25	0.52917 (19)	0.5717 (3)	-0.5084 (6)	0.0384 (10)
H25A	0.5227	0.5070	-0.5395	0.058*
H25B	0.4944	0.5992	-0.4806	0.058*
H25C	0.5550	0.5688	-0.4404	0.058*
C26	0.59983 (19)	0.5767 (3)	-0.6685 (5)	0.0329 (12)
H26A	0.5860	0.5136	-0.6923	0.049*
H26B	0.6300	0.5690	-0.6104	0.049*
H26C	0.6132	0.6106	-0.7405	0.049*
C27	0.70438 (16)	0.8849 (3)	-0.5894 (4)	0.0208 (9)
C28	0.70892 (19)	0.8820 (4)	-0.7288 (4)	0.0339 (11)
H28A	0.7468	0.8962	-0.7530	0.051*
H28B	0.6841	0.9297	-0.7641	0.051*
H28C	0.6987	0.8181	-0.7579	0.051*
C29	0.71837 (17)	0.9870 (3)	-0.5474 (4)	0.0252 (10)

H29A	0.7545	1.0052	-0.5788	0.038*
H29B	0.7189	0.9892	-0.4586	0.038*
H29C	0.6906	1.0316	-0.5781	0.038*
C30	0.74679 (15)	0.8160 (3)	-0.5365 (4)	0.0239 (11)
H30A	0.7837	0.8361	-0.5610	0.036*
H30B	0.7397	0.7512	-0.5667	0.036*
H30C	0.7442	0.8164	-0.4478	0.036*
C31	0.43785 (16)	0.9580 (3)	-0.2602 (4)	0.0197 (9)
H31A	0.4032	0.9923	-0.2783	0.024*
H31B	0.4582	0.9952	-0.1985	0.024*
C32	0.42489 (17)	0.8604 (3)	-0.2109 (4)	0.0196 (9)
H32A	0.4595	0.8281	-0.1876	0.024*
H32B	0.4020	0.8671	-0.1371	0.024*
C33	0.33508 (16)	0.8180 (3)	-0.2903 (4)	0.0207 (9)
H33A	0.3218	0.7915	-0.2133	0.031*
H33B	0.3158	0.7872	-0.3578	0.031*
H33C	0.3281	0.8871	-0.2922	0.031*
C34	0.40419 (19)	0.6974 (3)	-0.2756 (4)	0.0273 (10)
H34A	0.3912	0.6827	-0.1936	0.041*
H34B	0.4435	0.6830	-0.2816	0.041*
H34C	0.3839	0.6585	-0.3345	0.041*
C35	0.64330 (19)	0.2714 (3)	-0.3474 (4)	0.0322 (11)
H35	0.6526	0.2663	-0.2594	0.039*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Mo1	0.01303 (14)	0.01394 (14)	0.01511 (14)	-0.00271 (11)	-0.00087 (19)	-0.00026 (19)
Cl1	0.0472 (7)	0.0482 (7)	0.0413 (8)	-0.0196 (6)	0.0070 (6)	-0.0063 (6)
Cl2	0.0577 (9)	0.0264 (7)	0.1258 (15)	0.0056 (6)	0.0374 (10)	0.0150 (8)
Cl3	0.0442 (7)	0.0329 (6)	0.0368 (6)	0.0014 (5)	0.0031 (6)	0.0034 (6)
O1	0.0132 (12)	0.0155 (12)	0.0196 (17)	-0.0015 (9)	-0.0039 (12)	0.0029 (13)
O2	0.0140 (13)	0.0168 (14)	0.0287 (15)	-0.0036 (11)	0.0027 (12)	-0.0061 (12)
O3	0.0168 (13)	0.0163 (14)	0.0253 (14)	-0.0043 (11)	-0.0024 (12)	-0.0040 (12)
O4	0.0237 (16)	0.0274 (16)	0.0160 (14)	-0.0045 (13)	0.0041 (13)	0.0004 (13)
N1	0.0130 (16)	0.0124 (16)	0.0151 (16)	-0.0018 (13)	0.0032 (15)	-0.0010 (14)
N2	0.0199 (18)	0.0163 (17)	0.0150 (17)	-0.0020 (14)	0.0007 (15)	0.0021 (15)
C1	0.0195 (18)	0.0191 (18)	0.023 (2)	0.0011 (15)	0.000 (2)	0.004 (2)
C2	0.0219 (18)	0.0100 (17)	0.021 (3)	-0.0029 (13)	0.002 (2)	-0.005 (2)
C3	0.012 (2)	0.019 (2)	0.017 (2)	0.0016 (16)	0.0024 (17)	-0.0024 (18)
C4	0.017 (2)	0.023 (2)	0.016 (2)	0.0072 (17)	0.0020 (18)	0.0049 (18)
C5	0.026 (2)	0.015 (2)	0.019 (2)	0.0030 (17)	0.0015 (19)	0.0021 (18)
C6	0.0181 (18)	0.0177 (19)	0.025 (3)	-0.0039 (15)	-0.0005 (19)	0.0019 (19)
C7	0.020 (2)	0.015 (2)	0.024 (2)	0.0032 (16)	0.002 (2)	0.0037 (18)
C8	0.020 (2)	0.019 (2)	0.016 (2)	-0.0001 (17)	-0.0019 (18)	-0.0014 (18)
C9	0.017 (2)	0.026 (2)	0.027 (2)	-0.0010 (18)	-0.0070 (19)	0.001 (2)
C10	0.0185 (18)	0.0254 (19)	0.0278 (19)	0.0013 (15)	0.004 (3)	0.004 (3)
C11	0.019 (2)	0.024 (2)	0.021 (2)	0.0018 (18)	-0.0032 (18)	-0.0062 (19)

C12	0.028 (2)	0.013 (2)	0.034 (2)	0.0008 (17)	0.000 (2)	0.0013 (19)
C13	0.078 (4)	0.018 (2)	0.041 (3)	0.012 (2)	0.005 (3)	-0.001 (3)
C14	0.054 (3)	0.025 (3)	0.052 (3)	-0.006 (2)	-0.015 (3)	0.018 (2)
C15	0.036 (3)	0.024 (3)	0.047 (3)	-0.002 (2)	0.003 (2)	0.010 (2)
C16	0.016 (2)	0.019 (2)	0.022 (2)	-0.0004 (16)	-0.0020 (18)	-0.0059 (18)
C17	0.0076 (19)	0.020 (2)	0.023 (2)	0.0022 (16)	-0.0013 (17)	-0.0043 (19)
C18	0.0163 (19)	0.0179 (19)	0.019 (3)	0.0002 (15)	-0.0003 (18)	-0.0011 (18)
C19	0.016 (2)	0.022 (2)	0.0189 (19)	0.0057 (17)	-0.0060 (18)	-0.0044 (18)
C20	0.016 (2)	0.020 (2)	0.016 (2)	0.0086 (17)	-0.0001 (17)	-0.0037 (18)
C21	0.014 (2)	0.021 (2)	0.0174 (19)	0.0063 (17)	-0.0048 (16)	0.0043 (16)
C22	0.0174 (19)	0.0164 (19)	0.021 (3)	0.0004 (16)	-0.0033 (16)	-0.0059 (16)
C23	0.022 (2)	0.017 (2)	0.026 (2)	0.0048 (17)	-0.008 (2)	-0.0032 (19)
C24	0.034 (3)	0.033 (3)	0.037 (3)	0.013 (2)	-0.016 (2)	-0.019 (2)
C25	0.053 (3)	0.019 (2)	0.043 (3)	-0.0036 (18)	0.003 (4)	-0.012 (3)
C26	0.032 (3)	0.020 (2)	0.047 (3)	0.007 (2)	-0.010 (2)	-0.017 (2)
C27	0.014 (2)	0.028 (2)	0.020 (2)	0.0030 (17)	0.0020 (18)	0.0025 (19)
C28	0.026 (2)	0.052 (3)	0.024 (3)	0.001 (2)	0.007 (2)	0.004 (2)
C29	0.018 (2)	0.026 (2)	0.031 (2)	0.0006 (17)	0.0062 (18)	0.0032 (19)
C30	0.0125 (19)	0.028 (2)	0.031 (3)	0.0056 (16)	0.0024 (18)	-0.0025 (19)
C31	0.017 (2)	0.021 (2)	0.021 (2)	-0.0024 (17)	0.0019 (18)	-0.0041 (18)
C32	0.019 (2)	0.025 (2)	0.014 (2)	-0.0050 (18)	0.0008 (17)	0.0066 (18)
C33	0.015 (2)	0.025 (2)	0.022 (2)	-0.0053 (17)	0.0033 (18)	0.0001 (19)
C34	0.037 (3)	0.017 (2)	0.028 (2)	0.001 (2)	-0.003 (2)	0.009 (2)
C35	0.050 (3)	0.021 (2)	0.026 (2)	-0.005 (2)	-0.001 (2)	0.000 (2)

Geometric parameters (\AA , $^{\circ}$)

Mo1—O3	1.702 (2)	C15—H15B	0.9800
Mo1—O4	1.702 (3)	C15—H15C	0.9800
Mo1—O1	1.941 (2)	C16—C17	1.512 (5)
Mo1—O2	1.954 (3)	C16—H16A	0.9900
Mo1—N1	2.392 (3)	C16—H16B	0.9900
Mo1—N2	2.422 (3)	C17—C18	1.391 (5)
C11—C35	1.737 (4)	C17—C22	1.401 (5)
C12—C35	1.746 (4)	C18—C19	1.413 (6)
C13—C35	1.761 (5)	C19—C20	1.387 (5)
O1—C2	1.360 (4)	C19—C23	1.540 (6)
O2—C18	1.373 (4)	C20—C21	1.400 (5)
N1—C7	1.482 (5)	C20—H20	0.9500
N1—C31	1.492 (5)	C21—C22	1.388 (5)
N1—C16	1.517 (5)	C21—C27	1.536 (5)
N2—C34	1.483 (5)	C22—H22	0.9500
N2—C33	1.486 (5)	C23—C24	1.525 (6)
N2—C32	1.495 (5)	C23—C26	1.535 (6)
C1—C6	1.395 (5)	C23—C25	1.538 (7)
C1—C2	1.396 (5)	C24—H24A	0.9800
C1—C7	1.528 (6)	C24—H24B	0.9800
C2—C3	1.416 (5)	C24—H24C	0.9800

C3—C4	1.384 (5)	C25—H25A	0.9800
C3—C8	1.536 (5)	C25—H25B	0.9800
C4—C5	1.393 (6)	C25—H25C	0.9800
C4—H4	0.9500	C26—H26A	0.9800
C5—C6	1.394 (6)	C26—H26B	0.9800
C5—C12	1.531 (5)	C26—H26C	0.9800
C6—H6	0.9500	C27—C30	1.528 (5)
C7—H7A	0.9900	C27—C29	1.537 (6)
C7—H7B	0.9900	C27—C28	1.541 (6)
C8—C9	1.531 (5)	C28—H28A	0.9800
C8—C10	1.536 (6)	C28—H28B	0.9800
C8—C11	1.539 (5)	C28—H28C	0.9800
C9—H9A	0.9800	C29—H29A	0.9800
C9—H9B	0.9800	C29—H29B	0.9800
C9—H9BC	0.9800	C29—H29C	0.9800
C10—H10A	0.9800	C30—H30A	0.9800
C10—H10B	0.9800	C30—H30B	0.9800
C10—H10C	0.9800	C30—H30C	0.9800
C11—H11A	0.9800	C31—C32	1.501 (6)
C11—H11B	0.9800	C31—H31A	0.9900
C11—H11C	0.9800	C31—H31B	0.9900
C12—C13	1.519 (7)	C32—H32A	0.9900
C12—C14	1.528 (7)	C32—H32B	0.9900
C12—C15	1.534 (7)	C33—H33A	0.9800
C13—H13A	0.9800	C33—H33B	0.9800
C13—H13B	0.9800	C33—H33C	0.9800
C13—H13C	0.9800	C34—H34A	0.9800
C14—H14A	0.9800	C34—H34B	0.9800
C14—H14B	0.9800	C34—H34C	0.9800
C14—H14C	0.9800	C35—H35	1.0000
C15—H15A	0.9800		
O3—Mo1—O4	108.33 (13)	C17—C16—H16A	108.7
O3—Mo1—O1	98.81 (11)	N1—C16—H16A	108.7
O4—Mo1—O1	96.06 (12)	C17—C16—H16B	108.7
O3—Mo1—O2	99.29 (11)	N1—C16—H16B	108.7
O4—Mo1—O2	95.31 (12)	H16A—C16—H16B	107.6
O1—Mo1—O2	154.36 (12)	C18—C17—C22	119.4 (3)
O3—Mo1—N1	161.44 (12)	C18—C17—C16	118.4 (3)
O4—Mo1—N1	90.18 (12)	C22—C17—C16	121.6 (3)
O1—Mo1—N1	77.33 (10)	O2—C18—C17	117.3 (3)
O2—Mo1—N1	79.73 (10)	O2—C18—C19	120.7 (3)
O3—Mo1—N2	86.91 (12)	C17—C18—C19	121.9 (3)
O4—Mo1—N2	164.76 (12)	C20—C19—C18	116.0 (4)
O1—Mo1—N2	80.96 (11)	C20—C19—C23	122.0 (3)
O2—Mo1—N2	82.04 (11)	C18—C19—C23	122.0 (3)
N1—Mo1—N2	74.57 (10)	C19—C20—C21	124.0 (4)
C2—O1—Mo1	128.0 (2)	C19—C20—H20	118.0

C18—O2—Mo1	120.8 (2)	C21—C20—H20	118.0
C7—N1—C31	110.1 (3)	C22—C21—C20	117.9 (4)
C7—N1—C16	107.0 (3)	C22—C21—C27	123.1 (4)
C31—N1—C16	108.4 (3)	C20—C21—C27	118.9 (3)
C7—N1—Mo1	109.3 (2)	C21—C22—C17	120.6 (4)
C31—N1—Mo1	107.3 (2)	C21—C22—H22	119.7
C16—N1—Mo1	114.6 (2)	C17—C22—H22	119.7
C34—N2—C33	107.0 (3)	C24—C23—C26	107.7 (4)
C34—N2—C32	110.0 (3)	C24—C23—C25	109.5 (4)
C33—N2—C32	109.1 (3)	C26—C23—C25	106.8 (3)
C34—N2—Mo1	110.2 (2)	C24—C23—C19	109.7 (3)
C33—N2—Mo1	110.8 (2)	C26—C23—C19	111.2 (3)
C32—N2—Mo1	109.6 (2)	C25—C23—C19	111.9 (4)
C6—C1—C2	119.4 (4)	C23—C24—H24A	109.5
C6—C1—C7	115.5 (3)	C23—C24—H24B	109.5
C2—C1—C7	125.0 (3)	H24A—C24—H24B	109.5
O1—C2—C1	120.5 (3)	C23—C24—H24C	109.5
O1—C2—C3	118.8 (3)	H24A—C24—H24C	109.5
C1—C2—C3	120.7 (3)	H24B—C24—H24C	109.5
C4—C3—C2	116.8 (3)	C23—C25—H25A	109.5
C4—C3—C8	121.6 (3)	C23—C25—H25B	109.5
C2—C3—C8	121.6 (3)	H25A—C25—H25B	109.5
C3—C4—C5	124.6 (4)	C23—C25—H25C	109.5
C3—C4—H4	117.7	H25A—C25—H25C	109.5
C5—C4—H4	117.7	H25B—C25—H25C	109.5
C4—C5—C6	116.5 (4)	C23—C26—H26A	109.5
C4—C5—C12	122.4 (4)	C23—C26—H26B	109.5
C6—C5—C12	121.1 (4)	H26A—C26—H26B	109.5
C5—C6—C1	121.9 (4)	C23—C26—H26C	109.5
C5—C6—H6	119.1	H26A—C26—H26C	109.5
C1—C6—H6	119.1	H26B—C26—H26C	109.5
N1—C7—C1	118.5 (3)	C30—C27—C21	109.6 (3)
N1—C7—H7A	107.7	C30—C27—C29	108.7 (3)
C1—C7—H7A	107.7	C21—C27—C29	112.2 (3)
N1—C7—H7B	107.7	C30—C27—C28	108.4 (3)
C1—C7—H7B	107.7	C21—C27—C28	109.9 (3)
H7A—C7—H7B	107.1	C29—C27—C28	108.0 (4)
C9—C8—C10	106.8 (3)	C27—C28—H28A	109.5
C9—C8—C3	111.9 (3)	C27—C28—H28B	109.5
C10—C8—C3	110.8 (3)	H28A—C28—H28B	109.5
C9—C8—C11	107.1 (3)	C27—C28—H28C	109.5
C10—C8—C11	109.3 (3)	H28A—C28—H28C	109.5
C3—C8—C11	110.7 (3)	H28B—C28—H28C	109.5
C8—C9—H9A	109.5	C27—C29—H29A	109.5
C8—C9—H9B	109.5	C27—C29—H29B	109.5
H9A—C9—H9B	109.5	H29A—C29—H29B	109.5
C8—C9—H9BC	109.5	C27—C29—H29C	109.5
H9A—C9—H9BC	109.5	H29A—C29—H29C	109.5

H9B—C9—H9BC	109.5	H29B—C29—H29C	109.5
C8—C10—H10A	109.5	C27—C30—H30A	109.5
C8—C10—H10B	109.5	C27—C30—H30B	109.5
H10A—C10—H10B	109.5	H30A—C30—H30B	109.5
C8—C10—H10C	109.5	C27—C30—H30C	109.5
H10A—C10—H10C	109.5	H30A—C30—H30C	109.5
H10B—C10—H10C	109.5	H30B—C30—H30C	109.5
C8—C11—H11A	109.5	N1—C31—C32	110.8 (3)
C8—C11—H11B	109.5	N1—C31—H31A	109.5
H11A—C11—H11B	109.5	C32—C31—H31A	109.5
C8—C11—H11C	109.5	N1—C31—H31B	109.5
H11A—C11—H11C	109.5	C32—C31—H31B	109.5
H11B—C11—H11C	109.5	H31A—C31—H31B	108.1
C13—C12—C14	109.3 (4)	N2—C32—C31	111.7 (3)
C13—C12—C5	109.2 (4)	N2—C32—H32A	109.3
C14—C12—C5	112.0 (4)	C31—C32—H32A	109.3
C13—C12—C15	109.0 (4)	N2—C32—H32B	109.3
C14—C12—C15	106.7 (4)	C31—C32—H32B	109.3
C5—C12—C15	110.5 (3)	H32A—C32—H32B	107.9
C12—C13—H13A	109.5	N2—C33—H33A	109.5
C12—C13—H13B	109.5	N2—C33—H33B	109.5
H13A—C13—H13B	109.5	H33A—C33—H33B	109.5
C12—C13—H13C	109.5	N2—C33—H33C	109.5
H13A—C13—H13C	109.5	H33A—C33—H33C	109.5
H13B—C13—H13C	109.5	H33B—C33—H33C	109.5
C12—C14—H14A	109.5	N2—C34—H34A	109.5
C12—C14—H14B	109.5	N2—C34—H34B	109.5
H14A—C14—H14B	109.5	H34A—C34—H34B	109.5
C12—C14—H14C	109.5	N2—C34—H34C	109.5
H14A—C14—H14C	109.5	H34A—C34—H34C	109.5
H14B—C14—H14C	109.5	H34B—C34—H34C	109.5
C12—C15—H15A	109.5	Cl1—C35—Cl2	111.3 (3)
C12—C15—H15B	109.5	Cl1—C35—Cl3	110.1 (2)
H15A—C15—H15B	109.5	Cl2—C35—Cl3	110.0 (2)
C12—C15—H15C	109.5	Cl1—C35—H35	108.5
H15A—C15—H15C	109.5	Cl2—C35—H35	108.5
H15B—C15—H15C	109.5	Cl3—C35—H35	108.5
C17—C16—N1	114.4 (3)		
O3—Mo1—O1—C2	-136.2 (3)	C16—N1—C7—C1	172.9 (4)
O4—Mo1—O1—C2	-26.5 (3)	Mo1—N1—C7—C1	48.2 (4)
O2—Mo1—O1—C2	89.4 (4)	C6—C1—C7—N1	167.6 (4)
N1—Mo1—O1—C2	62.3 (3)	C2—C1—C7—N1	-15.7 (7)
N2—Mo1—O1—C2	138.4 (3)	C4—C3—C8—C9	0.4 (5)
O3—Mo1—O2—C18	133.7 (3)	C2—C3—C8—C9	-177.7 (4)
O4—Mo1—O2—C18	24.1 (3)	C4—C3—C8—C10	-118.6 (4)
O1—Mo1—O2—C18	-91.9 (3)	C2—C3—C8—C10	63.2 (5)
N1—Mo1—O2—C18	-65.1 (3)	C4—C3—C8—C11	119.9 (4)

N2—Mo1—O2—C18	-140.8 (3)	C2—C3—C8—C11	-58.3 (5)
O3—Mo1—N1—C7	-138.1 (3)	C4—C5—C12—C13	102.5 (5)
O4—Mo1—N1—C7	37.9 (2)	C6—C5—C12—C13	-76.6 (5)
O1—Mo1—N1—C7	-58.3 (2)	C4—C5—C12—C14	-18.7 (6)
O2—Mo1—N1—C7	133.3 (2)	C6—C5—C12—C14	162.1 (4)
N2—Mo1—N1—C7	-142.2 (2)	C4—C5—C12—C15	-137.6 (4)
O3—Mo1—N1—C31	-18.7 (5)	C6—C5—C12—C15	43.3 (6)
O4—Mo1—N1—C31	157.3 (2)	C7—N1—C16—C17	-85.1 (4)
O1—Mo1—N1—C31	61.2 (2)	C31—N1—C16—C17	156.2 (3)
O2—Mo1—N1—C31	-107.3 (2)	Mo1—N1—C16—C17	36.3 (4)
N2—Mo1—N1—C31	-22.8 (2)	N1—C16—C17—C18	-60.3 (5)
O3—Mo1—N1—C16	101.8 (4)	N1—C16—C17—C22	128.1 (4)
O4—Mo1—N1—C16	-82.2 (3)	Mo1—O2—C18—C17	64.7 (4)
O1—Mo1—N1—C16	-178.4 (3)	Mo1—O2—C18—C19	-114.9 (3)
O2—Mo1—N1—C16	13.2 (2)	C22—C17—C18—O2	-176.6 (3)
N2—Mo1—N1—C16	97.7 (3)	C16—C17—C18—O2	11.6 (5)
O3—Mo1—N2—C34	54.6 (3)	C22—C17—C18—C19	3.0 (6)
O4—Mo1—N2—C34	-126.1 (5)	C16—C17—C18—C19	-168.8 (4)
O1—Mo1—N2—C34	154.0 (3)	O2—C18—C19—C20	177.7 (3)
O2—Mo1—N2—C34	-45.2 (3)	C17—C18—C19—C20	-1.9 (6)
N1—Mo1—N2—C34	-126.7 (3)	O2—C18—C19—C23	-2.1 (6)
O3—Mo1—N2—C33	-63.7 (2)	C17—C18—C19—C23	178.4 (4)
O4—Mo1—N2—C33	115.6 (5)	C18—C19—C20—C21	-1.2 (6)
O1—Mo1—N2—C33	35.8 (2)	C23—C19—C20—C21	178.5 (4)
O2—Mo1—N2—C33	-163.5 (2)	C19—C20—C21—C22	3.1 (6)
N1—Mo1—N2—C33	115.0 (2)	C19—C20—C21—C27	-179.9 (4)
O3—Mo1—N2—C32	175.9 (3)	C20—C21—C22—C17	-2.0 (6)
O4—Mo1—N2—C32	-4.9 (6)	C27—C21—C22—C17	-178.8 (4)
O1—Mo1—N2—C32	-84.7 (2)	C18—C17—C22—C21	-1.0 (6)
O2—Mo1—N2—C32	76.0 (2)	C16—C17—C22—C21	170.6 (4)
N1—Mo1—N2—C32	-5.4 (2)	C20—C19—C23—C24	-106.9 (5)
Mo1—O1—C2—C1	-46.0 (6)	C18—C19—C23—C24	72.9 (5)
Mo1—O1—C2—C3	133.2 (3)	C20—C19—C23—C26	12.1 (5)
C6—C1—C2—O1	-179.0 (4)	C18—C19—C23—C26	-168.2 (4)
C7—C1—C2—O1	4.5 (7)	C20—C19—C23—C25	131.4 (4)
C6—C1—C2—C3	1.9 (7)	C18—C19—C23—C25	-48.9 (5)
C7—C1—C2—C3	-174.7 (4)	C22—C21—C27—C30	112.4 (4)
O1—C2—C3—C4	-179.4 (4)	C20—C21—C27—C30	-64.4 (5)
C1—C2—C3—C4	-0.2 (6)	C22—C21—C27—C29	-8.4 (5)
O1—C2—C3—C8	-1.2 (6)	C20—C21—C27—C29	174.7 (3)
C1—C2—C3—C8	178.0 (4)	C22—C21—C27—C28	-128.7 (4)
C2—C3—C4—C5	-0.5 (6)	C20—C21—C27—C28	54.5 (5)
C8—C3—C4—C5	-178.8 (4)	C7—N1—C31—C32	168.7 (3)
C3—C4—C5—C6	-0.4 (6)	C16—N1—C31—C32	-74.5 (4)
C3—C4—C5—C12	-179.5 (4)	Mo1—N1—C31—C32	49.8 (3)
C4—C5—C6—C1	2.1 (6)	C34—N2—C32—C31	155.2 (3)
C12—C5—C6—C1	-178.7 (4)	C33—N2—C32—C31	-87.7 (4)
C2—C1—C6—C5	-2.9 (7)	Mo1—N2—C32—C31	33.8 (4)

C7—C1—C6—C5 C31—N1—C7—C1	174.0 (4) −69.4 (5)	N1—C31—C32—N2	−58.0 (4)
-----------------------------	------------------------	---------------	-----------
