

(Dimethyl sulfoxide- κ O)[3-hydroxy-2-hydroxymethyl-2-(3-methoxy-2-oxido-benzylideneamino- κ^2 O²,N)propanolato- κ O]dioxomolybdenum(VI). Corrigendum**Yan Sui, Xiao-Niu Fang,* Qiu-Yan Luo, Hong-Mei Chen and Meng-Qiang Zhou**

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Key indicators: single-crystal X-ray study; $T = 295$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; disorder in main residue; R factor = 0.022; wR factor = 0.064; data-to-parameter ratio = 12.7.

The crystal structure of the title compound in the paper by Sui, Fang, Luo, Chen & Zhou [*Acta Cryst.* (2006), **E62**, m1994–m1996] has been re-refined to allow for identification of a disordered dimethyl sulfoxide ligand.

The structure reported by Sui *et al.* (2006) has been re-refined. The compound was originally determined by Rao *et al.* [*J. Chem. Soc. Dalton Trans.* (1998), 2383] and has been re-determined here to a significantly higher precision of the lattice parameters [$a = 14.3130$ (7) Å, $b = 9.2596$ (5) Å and $c = 14.8563$ (7) Å here *versus* $a = 14.305$ (3) Å, $b = 9.249$ (2) Å and $c = 14.860$ (3) Å reported by Rao *et al.*], bond lengths and s.u. values [*e.g.* Mo1–O6 = 1.6937 (17) Å here *versus* Mo1–O6 1.697 (4) Å reported by Rao *et al.*; $R = 0.022$ here *versus* $R = 0.050$ reported by Rao *et al.*]. The results of the current redetermination allow the identification of a disordered dimethyl sulfoxide ligand and a clarification of the nature of the intra- and intermolecular hydrogen bonding.

Experimental*Crystal data*

[Mo(C ₁₂ H ₁₅ NO ₅)O ₂ (C ₂ H ₆ OS)]	$V = 1779.74$ (15) Å ³
$M_r = 459.32$	$Z = 4$
Monoclinic, $P2_1/n$	Mo $K\alpha$ radiation
$a = 14.3130$ (7) Å	$\mu = 0.90$ mm ⁻¹
$b = 9.2596$ (5) Å	$T = 295$ K
$c = 14.8563$ (7) Å	$0.56 \times 0.39 \times 0.35$ mm
$\beta = 115.324$ (1)°	

Data collection

Bruker APEXII area-detector diffractometer	10668 measured reflections
Absorption correction: multi-scan (SADABS; Bruker, 2004)	3278 independent reflections
$T_{\min} = 0.632$, $T_{\max} = 0.746$	3056 reflections with $I > 2\sigma(I)$
	$R_{\text{int}} = 0.024$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.022$	40 restraints
$wR(F^2) = 0.064$	H-atom parameters constrained
$S = 1.07$	$\Delta\rho_{\text{max}} = 0.36$ e Å ⁻³
3278 reflections	$\Delta\rho_{\text{min}} = -0.44$ e Å ⁻³
259 parameters	

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
C1–H1 \cdots O5	0.93	2.22	2.874 (3)	127
C14–H14E \cdots O7 ⁱ	0.96	2.45	3.343 (4)	154
C8–H8A \cdots O6 ⁱⁱ	0.96	2.54	3.475 (4)	164
C11–H11A \cdots O7 ⁱⁱⁱ	0.97	2.57	3.534 (3)	170
O5–H5A \cdots O1 ⁱⁱⁱ	0.82	2.02	2.834 (2)	175
O5–H5A \cdots O2 ⁱⁱⁱ	0.82	2.53	2.970 (2)	115
O4–H4 \cdots O3 ^{iv}	0.82	1.99	2.801 (2)	171

Symmetry codes: (i) $-x + \frac{3}{2}, y - \frac{1}{2}, -z + \frac{5}{2}$; (ii) $x, y - 1, z$; (iii) $x - \frac{1}{2}, -y + \frac{3}{2}, z - \frac{1}{2}$; (iv) $-x + 1, -y + 2, -z + 2$.

Data collection: *APEX2* (Bruker, 2004); cell refinement: *APEX2* (Bruker, 2004); data reduction: *APEX2* (Bruker, 2004); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXL97* (Sheldrick, 2008); software used to prepare material for publication: *SHELXL97* (Sheldrick, 2008).

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: NG9064).

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