

Jin Lin, Hai-Bin Song and
Shan-Sheng Xu*State Key Laboratory of Elemento-Organic
Chemistry, College of Chemistry, Nankai
University, Tianjin 300071, People's Republic
of China

Correspondence e-mail: shanshengxu@126.com

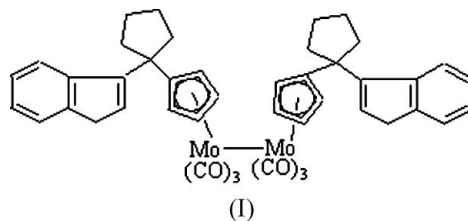
Key indicators

Single-crystal X-ray study
 $T = 293$ K
Mean $\sigma(\text{C}-\text{C}) = 0.007$ Å
 R factor = 0.037
 wR factor = 0.079
Data-to-parameter ratio = 15.7For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.Bis[[1-(1*H*-inden-3-yl)cyclopentyl]- η^5 -cyclopentadienyl]hexacarbonyldimolybdenum(I) (Mo—Mo)

The title centrosymmetric complex, $[\text{Mo}_2(\text{C}_{19}\text{H}_{19})(\text{CO})_6]$ or $\{[(\eta^5\text{-C}_5\text{H}_4)\text{C}(\text{CH}_2)_4(\text{C}_9\text{H}_7)]\text{Mo}(\text{CO})_3\}_2$, is a binuclear metal carbonyl complex containing an Mo—Mo single bond [3.2405 (9) Å]. The cyclopentane ring adopts an envelope conformation. The dihedral angle between the indene and cyclopentadiene rings is 57.0 (2)°.

Comment

Reaction of the lithium salt $[\text{C}_5\text{H}_4\text{-C}(\text{CH}_3)_2\text{-C}_9\text{H}_7]^- \cdot \text{Li}^+$ with $(\text{CH}_3\text{CN})_3\text{M}(\text{CO})_3$ ($M = \text{Mo}, \text{W}$) affords $\{[\text{C}_5\text{H}_4\text{-C}(\text{CH}_3)_2\text{-C}_9\text{H}_7]\text{M}(\text{CO})_3\}^- \cdot \text{Li}^+$, and subsequent reaction with MeI leads to the formation of a neutral complex $[\eta^5\text{-C}_5\text{H}_4\text{-C}(\text{CH}_3)_2\text{-C}_9\text{H}_7]\text{M}(\text{CO})_3\text{Me}$ (Alt *et al.*, 1993). Reaction of the dilithium salt $[\text{C}_5\text{H}_4\text{-C}(\text{CH}_3)_2\text{-C}_9\text{H}_6]^{2-} \cdot \text{Li}^{2+}$ with $\text{Mo}(\text{CO})_3(\text{CH}_3\text{CN})_3$ and oxidation with acetic acid and Fe^{III} produced a dimer, $\{\text{M}(\text{CO})_3[\eta^5\text{-C}_5\text{H}_4\text{-C}(\text{CH}_3)_2\text{-C}_9\text{H}_7]\}_2$, in which the indenyl ring was unmetallated (Fierro *et al.*, 1996). The title compound, $\{(\eta^5\text{-C}_5\text{H}_4)\text{C}(\text{CH}_2)_4(\text{C}_9\text{H}_7)\text{Mo}(\text{CO})_3\}_2$ (I), was obtained by the reaction of $[\text{C}_5\text{H}_5\text{-C}(\text{CH}_2)_4\text{-C}_9\text{H}_7]$ with $\text{Mo}(\text{CO})_3(\text{CH}_3\text{CN})_3$ in xylene.



A view of (I), with the atomic numbering scheme, is shown in Fig. 1, and selected bond lengths and angles are given in Table 1. Compound (I) is a centrosymmetric binuclear metal-carbonyl complex containing an Mo—Mo single bond. As shown in Fig. 1, only the cyclopentadienyl ring of the organic ligand coordinates with the Mo atom, and the indenyl group acts as a substituent. The Mo1...Cg1 distance is 2.0113 (19) Å, where Cg1 is the centroid of the cyclopentadiene ring. The Cg1—Mo1—C1, Cg1—Mo1—C2 and Cg1—Mo1—C3 angles are 116.09 (13), 125.76 (13) and 127.68 (13)°, respectively. The Mo—Mo bond distance is 3.2405 (9) Å and agrees with that observed in the analogous structures $(\eta^5\text{-C}_5\text{H}_5)_2\text{Mo}_2(\text{CO})_5(\text{CNCH}_3)$ [3.230 (1) Å; Adams *et al.*, 1973] and $(\eta^5\text{-C}_5\text{H}_5)_2\text{Mo}_2(\text{CO})_6$ [3.235 (1) Å; Adams *et al.*, 1974].

The two cyclopentadienyl rings are parallel by virtue of the centre of symmetry. The cyclopentane ring in the ligand adopts an envelope conformation, with atom C9 as the flap. The dihedral angle between the indene and cyclopentadiene rings is 57.0 (2)°. The crystal packing is stabilized by weak C—H...O interactions (Table 2).

Received 21 June 2006

Accepted 12 July 2006

Experimental

A solution of [C₅H₅-C(CH₂)₄-C₉H₇] (497 mg, 2 mmol) (Yan *et al.*, 1998) and Mo(CO)₃(CH₃CN)₃ [obtained by the reaction of Mo(CO)₆ (1000 mg, 3.79 mmol) with CH₃CN (15 ml) under refluxing for 4 h] (Tate *et al.*, 1962) in xylene (40 ml) was refluxed for 10 h. The solvent was removed under vacuum and the residue was chromatographed on an Al₂O₃ column using petroleum ether/CH₂Cl₂ (1:1) as eluent. The red band was collected and after several days red crystals were obtained (yield 0.06 g, 4%). Analysis calculated for C₄₄H₃₈Mo₂O₆: C 61.84, H 4.48%; found: C 61.78, H 4.65%.

Crystal data

[Mo₂(C₁₉H₁₉)(CO)₆] Z = 2
 M_r = 854.62 D_x = 1.556 Mg m⁻³
 Monoclinic, P2₁/c Mo Kα radiation
 a = 19.038 (4) Å μ = 0.74 mm⁻¹
 b = 7.7015 (16) Å T = 293 (2) K
 c = 13.091 (3) Å Plate, red
 β = 108.134 (3)° 0.30 × 0.20 × 0.08 mm
 V = 1824.1 (7) Å³

Data collection

Bruker SMART CCD 1000 area-detector diffractometer 9906 measured reflections
 φ and ω scans 3693 independent reflections
 Absorption correction: multi-scan 2385 reflections with I > 2σ(I)
 (SADABS; Sheldrick, 1996) R_{int} = 0.056
 T_{min} = 0.691, T_{max} = 1.000 θ_{max} = 26.3°
 (expected range = 0.652–0.943)

Refinement

Refinement on F² w = 1/[σ²(F_o²) + (0.0217P)² + 1.246P]
 R[F² > 2σ(F²)] = 0.037 where P = (F_o² + 2F_c²)/3
 wR(F²) = 0.079 (Δσ)_{max} = 0.001
 S = 1.01 Δρ_{max} = 0.52 e Å⁻³
 3693 reflections Δρ_{min} = -0.47 e Å⁻³
 235 parameters
 H-atom parameters constrained

Table 1

Selected geometric parameters (Å, °).

Mo1—C1	1.958 (4)	Mo1—C5	2.367 (4)
Mo1—C2	1.977 (4)	Mo1—C6	2.329 (4)
Mo1—C3	1.985 (4)	Mo1—C7	2.306 (4)
Mo1—C4	2.353 (4)	Mo1—C8	2.359 (4)
C1—Mo1—C2	77.65 (16)	C6—Mo1—C4	58.00 (15)
C1—Mo1—C3	78.33 (16)	C1—Mo1—C8	89.62 (15)
C2—Mo1—C3	106.13 (17)	C2—Mo1—C8	139.49 (15)
C1—Mo1—C7	90.54 (16)	C3—Mo1—C8	108.65 (15)
C2—Mo1—C7	105.28 (16)	C7—Mo1—C8	35.72 (13)
C3—Mo1—C7	143.38 (15)	C6—Mo1—C8	58.99 (14)
C1—Mo1—C6	122.14 (16)	C4—Mo1—C8	35.14 (13)
C2—Mo1—C6	95.71 (16)	C1—Mo1—C5	146.51 (15)
C3—Mo1—C6	153.31 (17)	C2—Mo1—C5	119.02 (17)
C7—Mo1—C6	35.24 (14)	C3—Mo1—C5	118.95 (17)
C1—Mo1—C4	120.81 (15)	C7—Mo1—C5	58.06 (15)
C2—Mo1—C4	152.75 (17)	C6—Mo1—C5	34.59 (15)
C3—Mo1—C4	97.73 (16)	C4—Mo1—C5	34.72 (14)
C7—Mo1—C4	58.36 (14)	C8—Mo1—C5	58.44 (14)

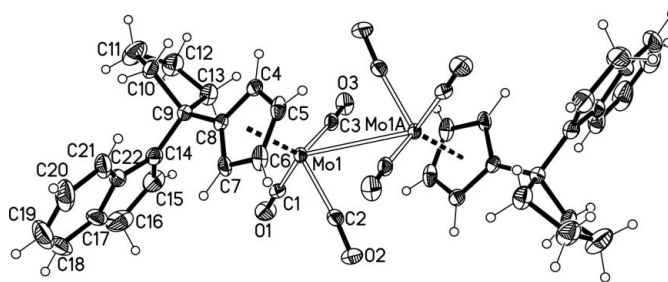


Figure 1

The structure of (I), showing 30% probability displacement ellipsoids. Atoms labelled with the suffix A and other unlabelled atoms are generated by the symmetry operation (−x, 1 − y, −z).

Table 2

Hydrogen-bond geometry (Å, °).

D—H···A	D—H	H···A	D···A	D—H···A
C5—H5···O3 ⁱ	0.98	2.40	3.334 (5)	160
C6—H6···O1 ⁱⁱ	0.98	2.53	3.499 (6)	171

Symmetry codes: (i) x, −y + ½, z − ½; (ii) x, −y + ½, z − ½.

H atoms were placed in calculated positions, with C—H = 0.93–0.98 Å, and refined using a riding model, with U_{iso}(H) = 1.2U_{eq}(parent atom).

Data collection: SMART (Bruker, 1998); cell refinement: SAINT (Bruker, 1999); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997a); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997a); molecular graphics: SHELXTL (Sheldrick, 1997b); software used to prepare material for publication: SHELXTL.

The authors thank the National Natural Science Foundation of China (grant Nos. 20472034 and 20574036), and the Research Fund for the Doctoral Program of Higher Education of China (grant No. 20050055008).

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