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Key indicators

Single-crystal X-ray study T = 100 KMean $\sigma(\text{O-C}) = 0.003 \text{ Å}$ R factor = 0.036 wR factor = 0.091Data-to-parameter ratio = 20.4

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

Redetermination of di- μ -bromo-bis[tetracarbonylmanganese(I)] at low temperature

The structure of the title compound, $[Mn_2Br_2(CO)_8]$, has previously been determined at room temperature using film data [Dahl & Wei (1963). *Acta Cryst.* **16**, 611–616]. The 100 K structure reported here is in good agreement with the previous study but of significantly higher precision.

Comment

Following the first synthesis of a transition metal carbonyl complex, considerable progress has been made towards extending this area of chemistry (Holleman-Wiberg, 1995). Earlier studies have shown that the carbonyl halides of manganese [Mn(CO)₅X] (X = Cl, Br, I) undergo thermal dissociation of coordinated carbon monoxide to give bis[halotetracarbonylmanganese(I)] (Brimm *et al.*, 1954). We now have found that storing manganese pentacarbonyl bromide [Mn(CO)₅Br] dissolved in CD₂Cl₂ *in vacuo* leads also to the formation of the the title dimeric manganese complex, [Mn(CO)₄Br]₂, (I), (Fig. 1).



The structure of (I) was first determined at room temperature using film data from Weissenberg exposures (Dahl & Wei, 1963). The present low-temperature determination yielded results of significantly improved precision (*e.g.*: the s.u.'s for the Mn-Br bonds decrease from 0.009 and 0.010 to 0.0005 Å, the s.u.'s for the Mn-C bonds decrease from 0.04



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and 0.05 to 0.003 Å). The improved precision helps to emphasize the difference between the Mn–C bond length *trans* to Br (mean value 1.887 Å) and *trans* to C (mean value 1.814 Å) (Table 1). The C–O bonds, on the other hand, show an inverse behaviour: those *trans* to Br (mean value 1.144 Å) are longer that those *trans* to C (mean value 1.134 Å). Both Mn atoms are octahedrally coordinated by four CO groups and two Br atoms bridging two Mn atoms. The molecule has chemical but not crystallographic D_{2h} symmetry.

Experimental

 $[Mn(CO)_5Br]$ (0.036 g, 0.13 mmol) and 1 ml CD_2Cl_2 were stored in a sealed and evacuated NMR tube for several days at rom temperature. Orange crystals of (I) suitable for X-ray diffraction were grown from this solution.

 $D_r = 2.502 \text{ Mg m}^{-3}$

Cell parameters from 20694

Mo Ka radiation

reflections

 $\theta=3.6{-}29.5^\circ$

 $\mu = 8.06~\mathrm{mm}^{-1}$

T = 100 (2) K

Block, orange

 $0.36 \times 0.20 \times 0.18 \text{ mm}$

Crystal data

 $[Mn_2Br_2(CO)_8]_2$ $M_r = 493.78$ Monoclinic, $P2_1/c$ a = 9.4545 (7) Å b = 11.5713 (8) Å c = 12.6818 (9) Å $\beta = 109.148$ (6)° V = 1310.64 (16) Å³ Z = 4

Data collection

STOE IPDS II two-circle	3694 independent reflections
diffractometer	3267 reflections with $I > 2\sigma(I)$
ω scans	$R_{\rm int} = 0.070$
Absorption correction: multi-scan	$\theta_{\rm max} = 29.7^{\circ}$
(MULABS; Spek, 2003; Blessing,	$h = -13 \rightarrow 13$
1995)	$k = -16 \rightarrow 16$
$T_{\min} = 0.160, \ T_{\max} = 0.225$	$l = -17 \rightarrow 17$
20694 measured reflections	

Refinement

-	
Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0606P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.036$	+ 0.3697P]
$wR(F^2) = 0.091$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.04	$(\Delta/\sigma)_{\rm max} = 0.001$
3694 reflections	$\Delta \rho_{\rm max} = 1.16 \text{ e } \text{\AA}^{-3}$
181 parameters	$\Delta \rho_{\rm min} = -1.38 \text{ e } \text{\AA}^{-3}$

Table 1

Selected geometric parameters (Å, °).

Mn1-C3	1.807 (3)	Mn2-C6	1.813 (3)
Mn1-C2	1.820 (3)	Mn2-C7	1.815 (3)
Mn1-C4	1.884 (3)	Mn2-C5	1.878 (3)
Mn1-C1	1.892 (3)	Mn2-C8	1.895 (3)
Mn1-Br2	2.5296 (5)	Mn2-Br1	2.5215 (5)
Mn1-Br1	2.5315 (5)	Mn2-Br2	2.5230 (5)
Mn2-Br1-Mn1	95.533 (16)	Mn2-Br2-Mn1	95.543 (17)

The highest peak and deepest hole in the final difference map are located 0.80 and 0.78 Å, respectively, from atom Br2.

Data collection: X-AREA (Stoe & Cie, 2001); cell refinement: X-AREA; data reduction: X-AREA; program(s) used to solve structure: SHELXS97 (Sheldrick, 1990); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: PLATON (Spek, 2003); software used to prepare material for publication: SHELXL97 and PLATON.

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