C31	-0.0708(4)	-0.4209 (4)	-0.2522 (4)	0.0520 (12)
C32	-0.1756 (6)	-0.4133 (5)	-0.2574 (6)	0.085 (2)
C33	-0.2432(7)	-0.4835 (6)	-0.2875 (8)	0.105 (3)
C 34	-0.2055 (7)	-0.5601 (6)	-0.3153 (7)	0.093 (3)
C 35	-0.1002(7)	-0.5666 (5)	-0.3109 (6)	0.090(2)
C 36	-0.0339(5)	-0.4990(5)	-0.2788(5)	0.073(2)

Table 2. Selected geometric parameters (Å, °)

	-		
Sn1-C13	2.118 (6)	C14—C15	1.394 (13)
Snl—Cì	2.120 (6)	C15C16	1.31 (2)
Snl—C7	2.134 (6)	C16C17	1.35 (2)
Sn1—O1	2.336 (4)	C17-C18	1.424 (15)
Sn1—Br1	2.6186(11)	C19-C24	1.393 (12)
P101	1.493 (4)	C19C20	1.416 (10)
P1-C31	1,789 (6)	C20-C21	1.349 (11)
P1	1 794 (6)	$C_{21} - C_{22}$	1.39(2)
P1C25	1.807 (6)	$C^{22} - C^{23}$	1.36(2)
C1C6	1.362 (10)	C23-C24	1 402 (13)
C_{1}	1 384 (9)	$C_{25} - C_{26}$	1 379 (9)
$C_1 - C_2$	1.351(11)	C25 C20	1 399 (10)
$C_2 = C_3$	1.331 (11)	C25-C37	1.380(11)
C4_C5	1.295(17)	C20_C27	1.368 (14)
C4C3	1.34 (2)	$C_{27} = C_{28}$	1.366 (14)
	1.430 (13)	$C_{20} - C_{29}$	1.300 (14)
C7	1.373(10)	C29-C30	1.348 (11)
C^{2}	1.379(10)	C31C32	1.307 (9)
	1.404 (12)	C31-C30	1.392 (9)
C9-C10	1.38 (2)	C32-C33	1.387 (11)
C10-C11	1.34 (2)	C33-C34	1.386 (13)
C11-C12	1.393 (11)	C34—C35	1.3/5(12)
C13C18	1.359 (10)	C35C36	1.355 (10)
C13C14	1.378 (10)		
C13-Sn1-C1	118.5 (2)	C18-C13-C14	119.3 (7)
C13—Sn1—C7	120.5 (2)	C18-C13-Sn1	120.8 (5)
C1—Sn1—C7	118.9 (2)	C14-C13-Sn1	119.8 (5)
C13-Sn1-O1	85.0 (2)	C13-C14-C15	119.8 (9)
C1—Sn1—O1	85.5 (2)	C16-C15-C14	120.2 (10)
C7—Sn1—O1	85.0 (2)	C15-C16-C17	122.7 (9)
C13—Sn1—Br1	94.1 (2)	C16-C17-C18	118.4 (10)
Cl = Sn1 = Br1	95.7 (2)	C13-C18-C17	119.6 (10)
C7Sn1Br1	94.7 (2)	C24-C19-C20	120.3 (7)
$\Omega_1 - Sn_1 - Br_1$	178 73 (11)	C24—C19—P1	121.1 (5)
01P1C31	111.3 (3)	C20-C19-P1	118.6 (6)
01 - P1 - C19	109.8 (3)	$C_{21} - C_{20} - C_{19}$	118.9 (9)
	107.6 (3)	C_{20} C_{21} C_{21} C_{22}	120.8 (9)
OI - PI - C25	1130(3)	C_{23} C_{23} C_{23} C_{23}	121.6 (8)
C31_P1_C25	108.1(3)	$C_{23} = C_{22} = C_{24}$	119.0 (10)
C19_P1_C25	106.8 (3)	$C_{22} = C_{23} = C_{24}$	119.4 (9)
$P_1 \cap P_2$	166 5 (3)	$C_{1}^{2} = C_{2}^{2} = C_{3}^{2}$	118 5 (6)
$C_{1} = 01 = 301$	116.2 (6)	C26-C25-C10	122 0 (5)
$C_{0} = C_{1} = C_{2}$	122 4 (5)	C20-C25-P1	119 5 (5)
$C_{0} = C_{1} = S_{01}$	122.4 (5)	$C_{30} = C_{25} = C_{25}$	119.9 (8)
$C_2 = C_1 = 311$	121.3(3)	$C_{21} = C_{20} = C_{23}$	110.7 (0)
$C_{3} = C_{2} = C_{1}$	121.4 (0)	$C_{20} = C_{27} = C_{20}$	171.3 (8)
$C_4 - C_3 - C_2$	123.4 (9)	$C_{29} = C_{20} = C_{27}$	110.2 (0)
	110.0 (9)	C_{30} C_{29} C_{28}	171.4 (9)
C4-C5-C6	120.3 (9)	$C_{29} = C_{30} = C_{23}$	121.4 (0)
$C_1 \rightarrow C_2 \rightarrow C_3$	119.9 (8)	$C_{22} - C_{21} - C_{20}$	119.2 (0)
(12 - (1 - 1))	118.0 (0)	C_{2} C_{2} C_{3} P_{1}	124.2 (3)
	121.9(5)	C30-C31-FI	124.3 (4)
C8-C/Snl	119.4 (5)	$C_{24} = C_{22} = C_{23}$	119.9(7)
U/U8U9	119.4 (9)	$C_{34} - C_{33} - C_{32}$	120.2 (7)
C10-C9-C8	120.5 (10)	C35-C34-C33	119.4 (7)
CII-CI0-C9	120.2 (8)	C30-C33-C34	120.2 (7)
C10-C11-C12	119.7 (9)	C35-C36-C31	121.1 (6)
C7C12C11	121.5 (8)		

H atoms were allowed to ride on their parent C atoms with $U(H) = 1.5U_{eq}$ of the parent atom.

Data collection: CAD-4 VAX/PC (Enraf-Nonius, 1988). Cell refinement: CAD-4 VAX/PC. Data reduction: Xtal3.0 (Hall & Stewart, 1990). Program(s) used to solve structure: SHELXS86 (Sheldrick, 1985). Program(s) used to refine structure: SHELXL93 (Sheldrick, 1993). Molecular graphics: ORTEPII (Johnson, 1976). Software used to prepare material for publication: SHELXL93.

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Lists of structure factors, anisotropic displacement parameters, Hatom coordinates and complete geometry have been deposited with the IUCr (Reference: JZ1056). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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$exo-\pi$ -(Benzonorbornadiene)tricarbonylchromium

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Abstract

The title compound, tricarbonyl[$(4a,5,6,7,8,8a-\eta)-1,4-di$ hydro-1.4-methanonaphthalene]chromium, $[Cr(C_{11}H_{10}) (CO)_{3}$, is a complex of chromium(0) in which the arene ring of the organic ligand is hexahapto-coordinated to the Cr atom. Three terminal carbonyl ligands form the triangular base of the molecule. The mode of coordination of the $Cr(CO)_3$ moiety to the ligand is such

that the free double bond of the benzonorbornadiene is oriented away from the Cr atom. The chromium– carbonyl tripod is rotated slightly from a perfectly staggered arrangement with respect to the arene ring and, hence, the molecule does not possess a plane of symmetry.

Comment

A large number of arenetricarbonylchromium compounds have been reported (Bruce, 1982). These compounds have been greatly studied, in part, because the Cr(CO)₃ unit alters the reactivity of substituted arenes in ways that depend on the stereochemistry of the complex (Bly & Maier, 1978). The prototype complex benzenetricarbonylchromium has a highly regular structure in which the Cr-CO bonds are staggered between the Cr-C bonds of the benzene ligand (Bailey & Dahl, 1965; Rees & Coppens, 1972). Complexes of substituted arenes are known in both staggered and eclipsed configurations, with the specific arrangement appearing to depend on the nature of the arene (Silverthorn, 1975). The structure of the most closely related compound to that reported here, $exo-\pi$ -(2-acetoxybenzonorbornadiene)tricarbonylchromium, has a staggered arrangement of the carbonyl ligands and an approximate plane of symmetry bisecting the Cr coordination sphere (Taylor, Griffith & Amma, 1976). The authors attribute the staggering of the carbonyl ligands to steric repulsions between the ligands and the bridgehead methylene group. The title benzonorbornadiene complex, (I) (Fig. 1), displays a similar structure in which the Cr atom is slightly displaced from the center of the arene ring, away from the norbornyl group. This is indicated by the shorter distances from the Cr atom to atoms C(9) and C(10) in comparison with those to C(2) and C(3) (Table 2).



The three carbonyl ligands form an almost regular triangular base for the complex. The two carbonyls adjacent to the norbornyl moiety are spread slightly apart so that the C(13)—Cr—C(14) angle is a few degrees larger than the angle that either C atom makes with C(12). The carbonyl tripod is not perfectly staggered between the benzene C atoms, but is rotated towards atoms C(9), C(2) and C(11). This displacement is evident in the bond angles between the carbonyl C atoms and those of the benzene ring, for example, C(13)—Cr—C(11) of 90.0 (3) and C(13)—Cr—C(3) of 93.9 (4). As a result, the molecule lacks a plane of symmetry that would otherwise bisect the benzonorbornadiene ligand,

the Cr atom and one of the carbonyl ligands. A study of the *exo* and *endo* isomers of tricarbonyl(2-methyl-1indanol)chromium found the tripod ideally staggered in the *endo* isomer and twisted in a very similar manner to the present compound in the *exo* isomer (Gentric, Le Borgne & Grandjean, 1978).



Fig. 1. ORTEPII diagram (Johnson, 1976) of the title compound showing the atomic labeling scheme. The displacement ellipsoids of the non-H atoms are shown at the 35% probability level.

The title compound was synthesized as part of a study of (alkenylarene)chromium compounds. Our original goal was to prepare the benzonorbornadiene complex with the free double bond directed towards the Cr atom in order to effect the coordination of the alkene *via* photolytic CO dissociation. The stereochemistry of the coordination of substituted indanes to chromium has been found to be highly dependent on reaction conditions, including solvent and temperature, with the ratio of *exo* and *endo* isomers varying widely for a given ligand (Gracey, Jackson, McMullen & Thompson, 1969). Our studies would seem to indicate the same is true of norbornylarenes.

Experimental

The benzonorbornadiene ligand was synthesized by the reaction of anthranilic acid, cyclopentadiene and *tert*-butyl nitrite in acetone and methylene chloride. The chromium complex was prepared by the reaction of hexacarbonylchromium (previously refluxed in acetonitrile) and the benzonorbornadiene ligand in refluxing glyme (1,2-dimethoxyethane). The product was isolated on a silica column with hexane and was recrystallized from a mixture of ether and pentane.

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Mo $K\alpha$ radiation $\lambda = 0.7107 \text{ Å}$

Cell parameters from 17 reflections $\theta = 9.2 - 13.3^{\circ}$ $\mu = 0.905 \text{ mm}^{-1}$ T = 296.2 KPlate

 $0.35 \times 0.10 \times 0.05$ mm

Yellow

Crystal data

$[Cr(C_{11}H_{10})(CO)_3]$
$M_{t} = 278.23$
Orthorhombic
$Pna2_1$
a = 15.225 (4) Å
<i>b</i> = 8.652 (3) Å
c = 9.306 (2) Å
$V = 1225(1) \text{ Å}^3$
Z = 4
$D_{\rm r} = 1.507 {\rm Mg} {\rm m}^{-3}$

Data collection

Rigaku AFC-6S diffractom-	$\theta_{\rm max} = 24.99^{\circ}$
eter	$h = 0 \rightarrow 18$
$\omega/2\theta$ scans	$k = 0 \rightarrow 11$
Absorption correction:	$l = 0 \rightarrow 10$
none	3 standard reflections
1281 measured reflections	monitored every 150
1281 independent reflections	reflections
755 observed reflections	intensity decay: 0.1%
$[I > 3\sigma(I)]$	

Refinement

Refinement on F	Weighting scheme based
R = 0.0328	on measured e.s.d.'s
wR = 0.0287	$(\Delta/\sigma)_{\rm max} = 0.0011$
S = 1.579	$\Delta \rho_{\rm max} = 0.19 \ {\rm e} \ {\rm \AA}^{-3}$
755 reflections	$\Delta \rho_{\rm min} = -0.29 \ {\rm e} \ {\rm \AA}^{-3}$
162 parameters	Atomic scattering factors
H atom parameters obtained	from International Tables
from difference map and	for X-ray Crystallography
not refined	(1974, Vol. IV, Table
	2.2B)

Table 1. Fractional atomic coordinates and equivalent isotropic displacement parameters ($Å^2$)

$U_{\text{eq}} = (1/3) \sum_{i} \sum_{i} U_{ii} a_i^* a_i^* \mathbf{a}_i . \mathbf{a}_i.$

	x	v	Z	U_{eq}
Cr(1)	0.83227 (6)	0.2955 (1)	0.4033	0.0457 (3)
O(1)	0.7313 (3)	0.5867 (5)	0.4504 (6)	0.076 (2)
O(2)	0.8863 (4)	0.4172 (7)	0.1154 (7)	0.094 (2)
O(3)	0.6707 (4)	0.1644 (7)	0.2711 (7)	0.086 (2)
C(1)	0.8557 (4)	-0.1010 (7)	0.408 (2)	0.063 (2)
C(2)	0.8666 (4)	0.0580 (8)	0.4784 (8)	0.046 (2)
C(3)	0.9427 (4)	0.1220 (7)	0.414 (1)	0.048 (2)
C(4)	0.9782 (5)	0.002(1)	0.3128 (10)	0.066 (3)
C(5)	1.0058 (5)	-0.1311 (8)	0.406 (2)	0.068 (2)
C(6)	0.9343 (5)	-0.1930 (8)	0.4649 (9)	0.067 (3)
C(7)	0.8891 (6)	-0.067(1)	0.2567 (9)	0.073 (3)
C(8)	0.8215 (5)	0.1322 (8)	0.5888 (8)	0.050(2)
C(9)	0.8540 (6)	0.276 (1)	0.6359 (9)	0.063 (3)
C(10)	0.9275 (6)	0.3422 (9)	0.578 (1)	0.073 (3)
C(11)	0.9738 (4)	0.2667 (9)	0.4656 (10)	0.063 (3)
C(12)	0.7712 (4)	0.4730 (8)	0.431 (1)	0.055 (2)
C(13)	0.8664 (6)	0.372 (1)	0.226 (1)	0.062 (3)
C(14)	0.7329 (5)	0.2155 (9)	0.3221 (8)	0.057 (2)

Table 2. Selected geometric parameters (Å, °)

Cr(1)—C(2)	2.233 (7)	C(1)—C(6)	1.533 (10)
Cr(1)—C(3)	2.256 (6)	C(1)C(7)	1.53 (1)
Cr(1)C(8)	2.237 (7)	C(2)—C(3)	1.417 (9)
Cr(1) - C(9)	2.196 (8)	C(2)—C(8)	1.392 (9)

		1.50(1)
2.246 (7)	C(3)—C(11)	1.422 (9)
1.814 (7)	C(4)—C(5)	1.50(1)
1.851 (9)	C(4)—C(7)	1.57 (1)
1.828 (7)	C(5)—C(6)	1.33 (1)
1.170 (7)	C(8)—C(9)	1.406 (10)
1.143 (10)	C(9)—C(10)	1.37(1)
1.148 (8)	C(10)—C(11)	1.42 (1)
1.533 (10)		
88.6 (3)	C(2)—C(3)—C(11)	118.3 (8)
93.9 (4)	C(4)—C(3)—C(11)	134.3 (7)
91.1 (3)	C(3)—C(4)—C(5)	105.6 (7)
90.1 (4)	C(3)—C(4)—C(7)	99.2 (6)
94.4 (3)	C(5)—C(4)—C(7)	98.1 (7)
90.0 (3)	C(4)—C(5)—C(6)	108.5 (7)
88.2 (4)	C(1)—C(6)—C(5)	106.7 (8)
87.4 (4)	C(1)—C(7)—C(4)	93.2 (6)
90.0 (4)	C(2)—C(8)—C(9)	117.6 (7)
103.5 (7)	C(8)—C(9)—C(10)	122.4 (8)
100.6 (7)	C(9)—C(10)—C(11)	120.2 (8)
99.2 (7)	C(3) - C(11) - C(10)	119.1 (7)
105.0 (7)	Cr(1) - C(12) - O(1)	179.3 (7)
132.6 (8)	Cr(1)—C(13)—O(2)	178.6 (9)
122.3 (7)	Cr(1)—C(14)—O(3)	179.5 (8)
106.9 (6)		
	$\begin{array}{c} 2.246 \ (7) \\ 1.814 \ (7) \\ 1.851 \ (9) \\ 1.828 \ (7) \\ 1.170 \ (7) \\ 1.143 \ (10) \\ 1.143 \ (10) \\ 1.148 \ (8) \\ 1.533 \ (10) \\ 88.6 \ (3) \\ 93.9 \ (4) \\ 91.1 \ (3) \\ 90.1 \ (4) \\ 90.1 \ (3) \\ 88.2 \ (4) \\ 90.0 \ (3) \\ 88.2 \ (4) \\ 87.4 \ (4) \\ 90.0 \ (4) \\ 103.5 \ (7) \\ 100.6 \ (7) \\ 99.2 \ (7) \\ 105.0 \ (7) \\ 132.6 \ (8) \\ 122.3 \ (7) \\ 106.9 \ (6) \end{array}$	$\begin{array}{cccccccccccccccccccccccccccccccccccc$

When the opposite enantiomer was refined under identical conditions, slightly poorer agreement indices were obtained: for observed data, R = 0.0337, wR = 0.0296 and S = 1.621.

Data collection: MSC/AFC Diffractometer Control Software (Molecular Structure Corporation, 1988). Cell refinement: MSC/AFC Diffractometer Control Software. Data reduction: TEXSAN (Molecular Structure Corporation, 1991). Program(s) used to solve structure: MITHRIL (Gilmore, 1983). Program(s) used to refine structure: TEXSAN. Software used to prepare material for publication: TEXSAN.

Lists of structure factors, anisotropic displacement parameters, Hatom coordinates, complete geometry and least-squares-planes data have been deposited with the IUCr (Reference: FG1100). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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