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# Tricarbonyl( $\eta^5$ -cyclopentadienyl)-(trifluorohydroxyborato)tungsten

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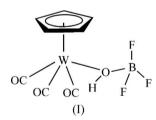
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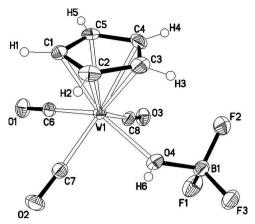
The title compound,  $[W(C_5H_5)(HOBF_3)(CO)_3]$ , has a fourlegged piano-stool geometry which is typically found for  $C_5H_5W(CO)_3X$  complexes. The HOBF<sub>3</sub><sup>-</sup> anion is the hydrolysis product of BF<sub>4</sub><sup>-</sup> and is coordinated *via* oxygen.

### Comment

 $[C_5H_5W(CO)_3]BF_4$  was first described by Beck *et al.* (1978) and used as a reactive intermediate (Werner *et al.*, 1987). All attempts to crystallize  $[(C_5H_5)W(CO)_3]BF_4$  for X-ray diffraction failed and the compound was described as extremely sensitive towards hydrolysis (Appel *et al.*, 1987; Beck *et al.*, 1989). We were able to obtain crystals of the first reaction product in the hydrolytic process for  $[(C_5H_5)W(CO)_3]BF_4$  that has been described before for the hydrolysis of  $[(C_5H_5)W-(CO)_3]BF_4$  to  $(CO)_5ReFBF_3$  (Raab *et al.*, 1981).

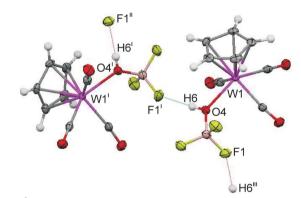


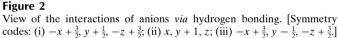
The crystal structure analysis showed that the HOBF<sub>3</sub> ligand is coordinated through the O atom (Fig. 1). The W–O distance of 2.155 (3) Å is significantly shorter than the W–O distance for coordinated water in  $(CO)_3(P^iPr_3)_2WOH_2$  of 2.320 (5) Å (Kubas *et al.*, 1992) and even shorter than the W–O distance for coordinated water in H<sub>2</sub>( $\eta^1$ -O<sub>2</sub>CBu<sup>t</sup>)<sub>2</sub>-(PMe<sub>3</sub>)<sub>3</sub>WOH<sub>2</sub> of 2.224 (2) Å (Zhu *et al.*, 2005). It is in the same range as the coordinated HOBF<sub>3</sub><sup>-</sup> anion in (CO)<sub>5</sub>ReO(H)BF<sub>3</sub> with an Re–O distance of 2.167 Å (Beck *et al.*, 1986). The HOBF<sub>3</sub> ligands of the title compound interact *via* an O–H···F hydrogen bond hydrogen bond which links the molecules into extended zigzag chains that run parallel to the [010] direction (Table 2 and Fig. 2).





The molecular structure of the title compound, with displacement ellipsoids drawn at the 50% probability level.





In an initial structure model, a weakly coordinating  $BF_4^$ anion was assumed rather than the hydrolysis product HOBF<sub>3</sub><sup>-</sup>. After convergence of the refinement at apparently convincing agreement factors, however, both the B-F distance and the displacement parameters (see Refinement section) indicated that the atom bridging W and B could be better refined as oxygen. When the bridging atom was modelled as an F atom, its distance from boron [1.485 (5) Å] was significantly longer than B-F distances found in the Cambridge Structural Database (Version 5.31; CSD; Allen, 2002) for tetrafluoroborate anions bound to metal centres via F [1.398 (23) Å] as observed for example in  $[CuL_2O(H)BF_3]$ - $BF_4$  (where L is 6-methyl-2,2'-bipyridine) containing both types of ligands discussed (Onggo et al., 1991). In the correct model, the B1–O4 distance [1.474 (5) Å] is nearly the same as the B-O distance in (CO)<sub>5</sub>ReO(H)BF<sub>3</sub> (1.476 Å) (Beck et al., 1986). IR spectra of the bulk material show a broad  $\nu(OH)$ band, indicating that the crystallizing product is only  $C_5H_5W(CO)_3O(H)BF_3$ .

For the cyclopentadienyl ligand, a slip distortion (distance of the W atom projection on the ring plane from the ring centre of gravity) of 0.106 Å is observed, with shorter metal-carbon bonds to C1 and C5, and longer W-C2, W-C3 and W-C4 distances (Table 1). The cyclopentadienyl (Cp) ligand

## metal-organic compounds

is tilted due to the strong *trans* effect of the CO ligands, as described previously for  $[Et_4N][C_5H_5Mo(CO)_2(\eta^2-O_2CO)]$  (Curtis & Han, 1985) and  $[C_5H_5TaCl_2(CO)_2(THF)]$  (THF is tetrahydrofuran) (Kwon & Curtis, 1990). This effect has also been explained theoretically (Chinn *et al.*, 1983).

### **Experimental**

The synthesis was carried out by the reaction of  $Cp(CO)_3WCH_3$  with HBF<sub>4</sub>·OEt<sub>2</sub> (1:2 molar ratio) in CH<sub>2</sub>Cl<sub>2</sub> for 30 min at 253 K. A deepred solution was formed immediately. The solvent was reduced at room temperature and red–orange crystals suitable for X-ray diffraction formed in 20% yield [m.p. 407 K (decomposition)]. IR [ATR (attenuated total reflection) technique using a diamond crystal, cm<sup>-1</sup>]: CO ligands: 2083 (*s*), 2017 (*s*), 1969 (*s*); HOBF<sub>3</sub><sup>-</sup> anion: 3401 (*br*), 1083 (*m*).

V = 1047.8 (3) Å<sup>3</sup>

Mo  $K\alpha$  radiation

0.14  $\times$  0.12  $\times$  0.06 mm

12253 measured reflections

2507 independent reflections

2289 reflections with  $I > 2\sigma(I)$ 

 $\mu = 11.07 \text{ mm}^-$ 

T = 100 K

 $R_{\rm int} = 0.025$ 

Z = 4

#### Crystal data

 $[W(C_{5}H_{3})(HOBF_{3})(CO)_{3}]$   $M_{r} = 417.79$ Monoclinic,  $P2_{1}/n$  a = 10.7679 (17) Å b = 8.5489 (14) Å c = 11.4313 (18) Å  $\beta = 95.297$  (2)°

#### Data collection

Bruker SMART APEX CCD diffractometer Absorption correction: multi-scan (SADABS; Bruker, 2000) T<sub>min</sub> = 0.345, T<sub>max</sub> = 0.437

#### Refinement

$R[F^2 > 2\sigma(F^2)] = 0.020$ wR(F^2) = 0.043	H atoms treated by a mixture of independent and constrained
S = 1.03	refinement
2507 reflections	$\Delta \rho_{\rm max} = 1.15 \text{ e} \text{ Å}^{-3}$
158 parameters	$\Delta \rho_{\rm min} = -0.74 \ {\rm e} \ {\rm \AA}^{-3}$
1	,

In an initial structure model, a weakly coordinated BF<sub>4</sub><sup>-</sup> anion was assumed rather than the hydrolysis product HOBF<sub>3</sub><sup>-</sup>. After convergence of the refinement at apparently convincing agreement factors, however, both the B1-F distance and displacement parameters show that atom X bridging W and B could be better refined as oxygen. In the initial tetrafluoroborate model, the B-X distance of 1.485 (5) Å was significantly longer than the B-F distances found in the CSD for tetrafluoroborate anions bonded to metals by F [metal-F-BF<sub>3</sub> = 1.398 (23) Å], and much longer than the terminal B-F distances [1.370(31) Å]. When X was refined as fluorine, a displacement parameter significantly larger than expected was obtained  $[U_{iso}(W1) =$ 0.01394 (6) Å<sup>2</sup>,  $U_{iso}(X) = 0.0374$  (7) Å<sup>2</sup> and  $U_{iso}(B1) = 0.0226$  (10) Å<sup>2</sup>], and two rigid-bond alerts were obtained in connection with these displacement parameters (Spek, 2009): PLAT230\_ALERT\_2\_C Hirshfeld Test Diff for F-B1 6.63 su and PLAT232\_ALERT\_2\_C Hirshfeld Test Diff (M-X) W1-F 9.52 su. According to the rigidbond postulate, the components of the anisotropic displacement parameters along chemical bonds are assumed to be equal in magnitude. Large differences supposedly indicate contamination of these parameters with other effects. Atomic sites assigned the wrong scattering type (e.g. Ag versus Br) should generate 'problem signals' with this test (Hirshfeld, 1976).

#### Table 1

Selected geometric parameters (Å, °).

W1-O4	2.155 (3)	W1-C3	2.363 (4)
W1-C5	2.271 (4)	O4-B1	1.474 (5)
W1-C1	2.274 (3)	F1-B1	1.397 (5)
W1-C2	2.323 (4)	F2-B1	1.380 (5)
W1-C4	2.347 (4)	F3-B1	1.376 (5)
B1-O4-W1	131.2 (2)	B1-O4-H6	120 (4)

Table	2
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Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$	
$O4{-}H6{\cdots}F1^i$	0.77 (5)	1.94 (5)	2.708 (4)	177 (6)	
Symmetry code: (i) $-x + \frac{3}{2}, y + \frac{1}{2}, -z + \frac{3}{2}$ .					

After correctly assigning the atomic scattering factor for O to X, the following results were obtained: (a) the displacement parameter for X became unexceptional  $[U_{iso}(X) = 0.0217 \text{ (6) } \text{Å}^2]$ ; (b) a local electron-density maximum, peak No. 5, represented the missing hydroxy H atom; free refinement of this atom resulted in reasonable geometry and an intermolecular hydrogen bond; (c) agreement factors R and wR2 improved (0.0201 versus 0.0208 and 0.043 versus 0.046, respectively); (d) the distance pattern became normal for HOBF<sub>3</sub> coordinated to metal (in the CSD, the average B–O distance is 1.472 Å); and (e) the above-mentioned rigid-bond alerts disappeared.

The Cp H atoms were refined in calculated positions, riding on C atoms, with C-H = 0.95 Å and  $U_{iso}(H) = 1.2U_{eq}(C)$ . The OH hydrogen was located in a difference Fourier map and refined freely.

Data collection: *SMART* (Bruker, 2000); cell refinement: *SAINT-Plus* (Bruker, 2000); data reduction: *SAINT-Plus*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *XP* in *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *XCIF* in *SHELXTL*.

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: EG3068). Services for accessing these data are described at the back of the journal.

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