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## Structure Reports

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## Key indicators

Single-crystal X-ray study
$T=110 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.003 \AA$
$R$ factor $=0.049$
$w R$ factor $=0.138$
Data-to-parameter ratio $=19.8$
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

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## 2-(Diphenylphosphinoylmethyl)pyridinium tetrafluoroborate

In the title compound, $\left[\mathrm{OPPh}_{2}\left(2-\mathrm{CH}_{2} \mathrm{C}_{5} \mathrm{H}_{4} \mathrm{NH}\right)\right] \mathrm{BF}_{4}$, the phosphorus atom displays a tetrahedral geometry, with the nitrogen of the pyridyl group protonated and having tetrafluoroborate as counter-anion.

## Comment

Vapor diffusion of diethyl ether into a solution reaction of $\mathrm{Cu}\left(\mathrm{BF}_{4}\right)_{2}$ with 2-pyridyldiphenylphosphine in acetonitrile at 278 K, provided two easily distinguished kinds of crystals: blue block-shaped crystals corresponding to the reaction product of the $\mathrm{Cu}^{\text {II }}$ salt with the bidentate phosphine ligand and colorless block-shaped crystals of the title compound, (I). An X-ray crystallographic investigation showed (I) to be the oxidized ligand, with further protonation of the nitrogen of the pyridyl group (Fig. 1) due to the presence of water in the system, the charge being balanced by $\mathrm{BF}_{4}{ }^{-}$present in the solution. The socalled hemilabile hybrid ligand (Minghetti et al., 1998) presents a very close to tetrahedral geometry, with the angles around the P atom ranging from 105.61 (8) to 112.31 (7) ${ }^{\circ}$ (Table 1). The bond lengths and angles are in normal ranges (Allen et al., 1987).

(I)

In the crystal structure, the molecules are linked into two chains running in different directions (Fig. 2) by $\mathrm{C}-\mathrm{H} \cdots \mathrm{F}$ and $\mathrm{N}-\mathrm{H} \cdots \mathrm{O}$ intermolecular hydrogen bonds (Table 2).

## Experimental

The title compound, (I), was obtained from the oxidation and protonation of 2-pyridyldiphenylphosphine from a reaction solution containing $\mathrm{Cu}\left(\mathrm{BF}_{4}\right)_{2} \quad(0.102 \mathrm{~g}, 0.4 \mathrm{mmol})$; the phosphine ligand ( $0.110 \mathrm{~g}, 0.4 \mathrm{mmol}$ ) was synthesized as reported in the literature with slight modifications of the procedure (Akermark et al., 1986). Diffraction-quality crystals of (I) were obtained by vapor diffusion of diethyl ether into a concentrated acetonitrile solution of the reaction mixture in the presence of air.


Figure 1
A view of the molecular structure of the cation of (I), with displacement ellipsoids drawn at the $50 \%$ probability level

## Crystal data

$\mathrm{C}_{18} \mathrm{H}_{17} \mathrm{NOP}^{+} \cdot \mathrm{BF}_{4}{ }^{-}$
$M_{r}=381.11$
Monoclinic, C2/c
$a=16.7528$ (15) $\AA$
$b=13.9287$ (17) $\AA$
$c=15.661$ (2) $\AA$
$\beta=95.314$ (4) ${ }^{\circ}$
$V=3638.7(7) \AA^{3}$

## Data collection

Bruker APEX X8 CCD areadetector diffractometer $\varphi$ and $\omega$ scans
Absorption correction: multi-scan (SADABS, Bruker, 2000)
$T_{\text {min }}=0.958, T_{\text {max }}=0.960$

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.049$
$w R\left(F^{2}\right)=0.138$
$S=1.04$
4726 reflections
239 parameters
H atoms treated by a mixture of independent and constrained refinement

$$
Z=8
$$

$D_{x}=1.391 \mathrm{Mg} \mathrm{m}^{-3}$
Mo $K \alpha$ radiation
$\mu=0.20 \mathrm{~mm}^{-1}$
$T=110$ (2) K
Block, colorless
$0.29 \times 0.25 \times 0.21 \mathrm{~mm}$

14958 measured reflections 4726 independent reflections 3915 reflections with $I>2 \sigma(I)$

$$
R_{\mathrm{int}}=0.041
$$

$$
\theta_{\max }=29.0^{\circ}
$$

$$
\begin{aligned}
& \begin{array}{l}
w=1 /\left[\sigma^{2}\left(F_{\mathrm{o}}^{2}\right)+(0.0684 P)^{2}\right. \\
\quad+4.7739 P] \\
\text { where } P=\left(F_{\mathrm{o}}^{2}+2 F_{\mathrm{c}}^{2}\right) / 3 \\
(\Delta / \sigma)_{\max }=0.001 \\
\Delta \rho_{\max }=0.65 \mathrm{e} \AA^{-3} \\
\Delta \rho_{\min }=
\end{array}-_{0.57 \mathrm{e}^{-3}}
\end{aligned}
$$

Table 1
Selected geometric parameters ( $\left({ }^{\circ},{ }^{\circ}\right)$.

| P1-O1 | $1.4939(12)$ | $\mathrm{P} 1-\mathrm{C} 13$ | $1.8000(17)$ |
| :--- | :--- | :--- | :--- |
| $\mathrm{P} 1-\mathrm{C} 7$ | $1.7946(17)$ | $\mathrm{P} 1-\mathrm{C} 1$ | $1.8283(18)$ |
|  |  |  |  |
| $\mathrm{O} 1-\mathrm{P} 1-\mathrm{C} 7$ | $112.31(7)$ | $\mathrm{O} 1-\mathrm{P} 1-\mathrm{C} 1$ | $111.51(8)$ |
| $\mathrm{O} 1-\mathrm{P} 1-\mathrm{C} 13$ | $110.04(8)$ | $\mathrm{C} 7-\mathrm{P} 1-\mathrm{C} 1$ | $105.61(8)$ |
| $\mathrm{C} 7-\mathrm{P} 1-\mathrm{C} 13$ | $108.97(8)$ | $\mathrm{C} 13-\mathrm{P} 1-\mathrm{C} 1$ | $108.22(8)$ |



Figure 2
Packing diagram of (I), showing the $\mathrm{C}-\mathrm{H} \cdots \mathrm{F}$ and $\mathrm{N}-\mathrm{H} . \cdots \mathrm{O}$ hydrogen bonds as dashed lines.

Table 2
Hydrogen-bond geometry ( $\AA,^{\circ}$ ).

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| $\mathrm{~N} 1-\mathrm{H} 1 C \cdots \mathrm{O}^{\mathrm{i}}$ | $0.96(3)$ | $1.67(3)$ | $2.6212(18)$ | $169(2)$ |
| $\mathrm{C} 1-\mathrm{H} 1 A \cdots \mathrm{~F}^{\mathrm{ii}}$ | 0.99 | 2.38 | $3.356(2)$ | 169 |
| $\mathrm{C} 18-\mathrm{H} 18 \cdots \mathrm{~F}^{\mathrm{ii}}$ | 0.95 | 2.49 | $3.265(3)$ | 139 |
| $\mathrm{C} 3-\mathrm{H} 3 \cdots \mathrm{F3}^{\mathrm{ii}}$ | 0.95 | 2.46 | $3.345(2)$ | 155 |
| $\mathrm{C} 6-\mathrm{H} 6 \cdots \mathrm{~F}^{\mathrm{iii}}$ | 0.95 | 2.40 | $3.271(3)$ | 152 |
| Symmetry codes: (i) $-x, y,-z+\frac{1}{2} ;$; (ii) $-x+\frac{1}{2}, y-\frac{1}{2},-z+\frac{1}{2} ;$ (iii) $-x, y-1,-z+\frac{1}{2}$ |  |  |  |  |

The H atom attached to the N atom was located in a difference map and refined isotropically. All other H atoms were positioned geometrically, with $\mathrm{C}-\mathrm{H}=0.95$ or $0.99 \AA$ for aromatic and methylene H atoms, respectively, and constrained to ride on their parent atoms, with $U_{\text {iso }}(\mathrm{H})=1.2 U_{\text {eq }}(\mathrm{C})$.

Data collection: APEX2 (Bruker, 2003); cell refinement: APEX2; data reduction: SAINT-Plus (Bruker, 2003); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Sheldrick, 2000); software used to prepare material for publication: SHELXTL.

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