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## Structure Reports

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## Key indicators

Single-crystal X-ray study
$T=150 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.003 \AA$
$R$ factor $=0.048$
$w R$ factor $=0.124$
Data-to-parameter ratio $=14.7$
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.
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## 2-(2,4-Dimethylphenyl)-4-isopropyl-1-methylisoquinolinium hexafluorophosphate

The title compound, $\mathrm{C}_{21} \mathrm{H}_{24} \mathrm{~N}^{+} \cdot \mathrm{PF}_{6}^{-}$, crystallizes in centrosymmetric hydrogen-bonded clusters consisting of two cations and two anions, via weak aromatic $\mathrm{C}-\mathrm{H} \cdots \mathrm{F}$ interactions. The isoquinolinium and dimethylphenyl moieties are not coplanar.

## Comment

In the context of the study of ring-closure reactions with imines (Diederen et al., 1998), the crystal structure of the title reaction product, (I), was determined. The title compound crystallizes in the triclinic space group $P \overline{1}$ and features hydrogen-bonded cation/anion pairs, which are related by the centre of symmetry. These pairs are joined by weak aromatic $\mathrm{C}-\mathrm{H} \cdots \mathrm{F}$ hydrogen bonds, between H5 and $\mathrm{F} 4^{i}$ [symmetry code: (i) $1-x, 1-y,-z ; 2.51 \AA$ A , and between H 19 and $\mathrm{F}^{\mathrm{ii}}$ [symmetry code: (ii) $1+x, y, z ; 2.53 \AA$. These hydrogen bonds are in the 2.1-2.6 $\AA$ region reported for $\mathrm{C}-\mathrm{H} \cdots \mathrm{F}(\mathrm{P})$ interactions (Grepioni et al., 1998), and are likely to be the reason that the $\mathrm{PF}_{6}{ }^{-}$anion is not disordered in this case.



The isoquinolinium ring system and the 2,4-dimethylphenyl group are not coplanar, probably because of unfavourable steric interactions between H 19 and the H atoms on C 10 , resulting in a dihedral angle of $71.58(8)^{\circ}$ between their leastsquares planes.

## Experimental

The compound was synthesized by J. Diederen (University of Amsterdam), and recrystallized from dichloromethane and pentane.

## Crystal data

| $\mathrm{C}_{21} \mathrm{H}_{24} \mathrm{~N}^{+} \cdot \mathrm{PF}_{6}{ }^{-}$ | $Z=2$ |
| :--- | :--- |
| $M_{r}=435.38$ | $D_{x}=1.438 \mathrm{Mg} \mathrm{m}$ |
| Triclinic, $P \overline{1} \overline{1}$ | Mo $K \alpha$ radiation |
| $a=8.7413(4) \AA$ | Cell parameters from 9057 |
| $b=9.0199(5) \AA$ | $\quad$ reflections |
| $c=12.8984(5) \AA$ | $\mu=1.6-26.0^{\circ}$ |
| $\alpha=93.027(3)^{\circ}$ | $T=0.20 \mathrm{~mm}^{-1}$ |
| $\beta=98.039(3)^{\circ}$ | Block, colourless |
| $\gamma=90.032(2)^{\circ}$ | $0.2 \times 0.14 \times 0.11 \mathrm{~mm}$ |
| $V=1005.55(8) \AA^{\circ}$ |  |

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Figure 1
View of the title compound, with the atom-numbering scheme. Displacement ellipsoids are drawn at the $50 \%$ probability level.

## Data collection

Nonius KappaCCD diffractometer $\varphi$ and $\omega$ scans
Absorption correction: multi-scan (MULABS; Blessing, 1995) $T_{\text {min }}=0.80, T_{\text {max }}=0.97$
9057 measured reflections 3931 independent reflections

2547 reflections with $I>2 \sigma(I)$
$R_{\text {int }}=0.063$
$\theta_{\text {max }}=26.0^{\circ}$
$h=-10 \rightarrow 10$
$k=-11 \rightarrow 9$
$l=-15 \rightarrow 15$

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.048$
$w R\left(F^{2}\right)=0.124$
$S=0.99$
3931 reflections
267 parameters

H -atom parameters constrained
$w=1 /\left[\sigma^{2}\left(F_{\mathrm{o}}{ }^{2}\right)+(0.0614 P)^{2}\right]$
where $P=\left(F_{\mathrm{o}}{ }^{2}+2 F_{\mathrm{c}}{ }^{2}\right) / 3$
$(\Delta / \sigma)_{\max }<0.001$ 。
$\Delta \rho_{\max }=0.21 \mathrm{e}_{\AA^{-3}}$
$\Delta \rho_{\min }=-0.39 \mathrm{e}^{-3}$

Table 1
Hydrogen-bond geometry ( $\AA^{\circ}{ }^{\circ}$ ).

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| C5-H5 $\cdots \mathrm{F}^{\mathrm{i}}$ | 0.95 | 2.51 | $3.445(2)$ | 167 |
| C19-H19 $\cdots \mathrm{F}^{\mathrm{ii}}$ | 0.95 | 2.53 | $3.384(3)$ | 149 |

Symmetry codes: (i) $-x+1,-y+1,-z$; (ii) $x+1, y, z$.
All H atoms were placed in geometrically idealized positions ( $\mathrm{C}-$ $\mathrm{H}=0.95-1.00 \AA$ ) and constrained to ride on their parent atoms, with $U_{\text {iso }}(\mathrm{H})=1.5 U_{\text {eq }}(\mathrm{C})$ for methyl H atoms and $U_{\text {iso }}(\mathrm{H})=1.2 U_{\text {eq }}(\mathrm{C})$ for all other H atoms.

Data collection: COLLECT (Hooft, 1998); cell refinement: HKL2000 (Otwinowski \& Minor, 1997); data reduction: HKL2000; program(s) used to solve structure: SIR97 (Altomare et al., 1999); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: PLATON (Spek, 2003); software used to prepare material for publication: PLATON.


Figure 2
Hydrogen-bonded (dashed lines) centrosymmetric cluster of two cations and two anions. [Symmetry codes: $(A) 1+x, y, z ;(B) 1-x, 1-y,-z ;(C)$ $2-x, 1-y,-z$.]


Figure 3
Packing diagram. View along the crystallographic $b$ axis.

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