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Key indicators

Single-crystal X-ray study
 T = 173 K
 Mean $\sigma(\text{C}-\text{C}) = 0.005 \text{ \AA}$
 R factor = 0.033
 wR factor = 0.075
 Data-to-parameter ratio = 6.9

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

Tetra-*n*-butylammonium tetrakis(pentafluorophenyl)borate

In the title compound, $\text{C}_{16}\text{H}_{36}\text{N}^+\cdot\text{C}_{24}\text{BF}_{20}^-$, the geometric parameters do not show any unusual values. The four *n*-butyl chains adopt an all-*trans* conformation.

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Comment

Very strong cationic Lewis acids are only available in the presence of weakly coordinating counter-ions such as $[\text{B}(\text{C}_6\text{F}_5)_4]^-$. These anions also play an important role in electrochemistry. The salt $[\text{nBu}_4\text{N}]^+\cdot[\text{B}(\text{C}_6\text{F}_5)_4]^-$ possesses a low reactivity towards reducing and oxidizing agents. Moreover, conductance measurements show that the dissociation constants of $[\text{nBu}_4\text{N}]^+\cdot[\text{B}(\text{C}_6\text{F}_5)_4]^-$ solutions in solvents with low polarity are greater than those of smaller traditional anions (LeSuer *et al.*, 2004). Therefore, we are convinced that tetra-*n*-butylammonium tetrakis(pentafluorophenyl)borate, (I), is an ideal supporting electrolyte. We have synthesized (I) and we report its crystal structure here.

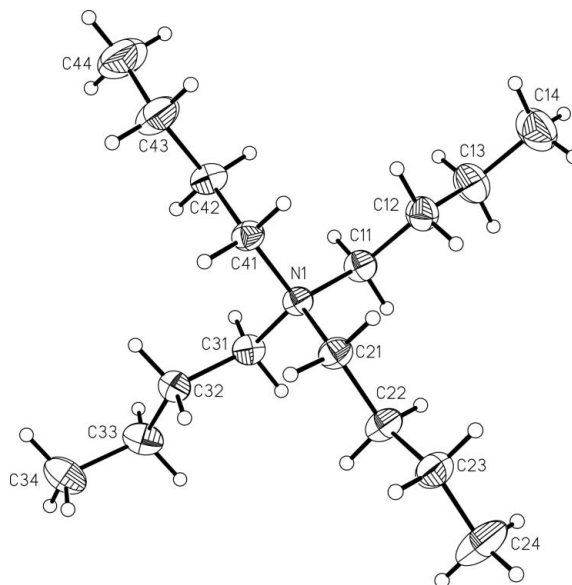
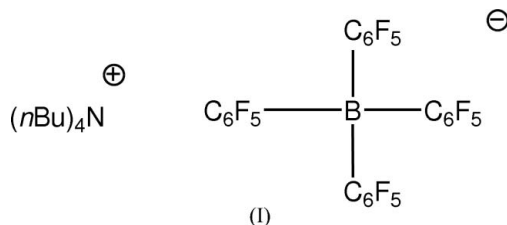


Figure 1
 Perspective view of the cation of (I), with the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level.

Perspective views of the cation and anion of (I) are shown in Figs. 1 and 2, respectively. Bond lengths and angles can be regarded as normal (Cambridge Crystallographic Database; Version 1.6 plus three updates; Mogul Version 1.0; Allen, 2002). The four *n*-butyl chains adopt an all-*trans* conformation.

Experimental

Compound (I) was synthesized according to the literature procedure of LeSuer *et al.* (2004). Colourless crystals of the title compound suitable for X-ray diffraction were grown from a methanol solution at 253 K.

Crystal data

$C_{16}H_{36}N^+ \cdot C_{24}BF_{20}^-$
 $M_r = 921.51$
 Monoclinic, *Cc*
 $a = 24.104$ (2) Å
 $b = 12.8925$ (12) Å
 $c = 17.5363$ (15) Å
 $\beta = 130.629$ (5)°
 $V = 4135.9$ (7) Å³
 $Z = 4$

$D_x = 1.480$ Mg m⁻³
 Mo $K\alpha$ radiation
 Cell parameters from 10062 reflections
 $\theta = 3.6$ – 25.6 °
 $\mu = 0.15$ mm⁻¹
 $T = 173$ (2) K
 Block, colourless
 $0.42 \times 0.37 \times 0.21$ mm

Data collection

Stoe IPDS-II two-circle diffractometer
 ω scans
 Absorption correction: none
 10176 measured reflections
 3854 independent reflections

3371 reflections with $I > 2\sigma(I)$
 $R_{int} = 0.035$
 $\theta_{max} = 25.6$ °
 $h = -29 \rightarrow 29$
 $k = -15 \rightarrow 14$
 $l = -21 \rightarrow 17$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.033$
 $wR(F^2) = 0.075$
 $S = 1.00$
 3854 reflections
 560 parameters
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0476P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{max} < 0.001$
 $\Delta\rho_{max} = 0.25$ e Å⁻³
 $\Delta\rho_{min} = -0.19$ e Å⁻³
 Extinction correction: *SHELXL97*
 Extinction coefficient: 0.0036 (4)

Table 1

Selected bond lengths (Å).

N1—C11	1.527 (4)	B1—C81	1.657 (4)
N1—C31	1.528 (4)	B1—C51	1.660 (5)
N1—C41	1.529 (3)	B1—C71	1.666 (4)
N1—C21	1.529 (3)	B1—C61	1.681 (4)

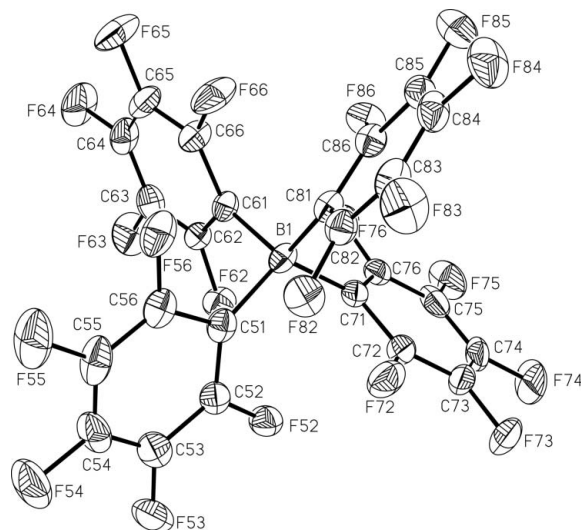


Figure 2

Perspective view of the anion of (I), with the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level.

H atoms were located in a difference map, but refined with fixed individual displacement parameters [$U_{iso}(H) = 1.2U_{eq}(C_{methylene})$ or $U_{iso}(H) = 1.5U_{eq}(C_{methyl})$] using a riding model, with C—H = 0.98 and 0.99 Å for C_{methyl} and $C_{methylene}$, respectively. In the absence of significant anomalous scatterers, Friedel pairs were merged.

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: *XP* in *SHELXTL-Plus* (Sheldrick, 1991); software used to prepare material for publication: *SHELXL97* and *PLATON* (Spek, 2003).

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