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## Key indicators

Single-crystal X-ray study
$T=173 \mathrm{~K}$
Mean $\sigma(\mathrm{N}-\mathrm{C})=0.008 \AA$
Disorder in main residue
$R$ factor $=0.033$
$\omega R$ factor $=0.072$
Data-to-parameter ratio $=21.5$
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.
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# Dimethylethylammonium bromide 

The title compound, $\mathrm{C}_{4} \mathrm{H}_{12} \mathrm{~N}^{+} \cdot \mathrm{Br}^{-}$, is a hydrolysis product of the 1-(dibromoboryl)ferrocene-dimethylethylamine (1/1) adduct. The metric symmetry is apparently orthorhombic $C$ centred, but the intensity statistics indicate unambigously the correct Laue symmetry, viz. primitive monoclinic. The ethyl group is disordered over two sites. In addition to the hydrogen bond from the NH group to the $\mathrm{Br}^{-}$ion, the packing is stabilized by several weak $\mathrm{C}-\mathrm{H} \cdots \mathrm{Br}$ hydrogen bonds.

## Comment

Recently, our group reported the properties and X-ray crystal structure determination of triferrocenylboroxine (Ma et al., 2002). We report here the hydrolysis of the 1-(dibromo-boryl)ferrocene-dimethylethylamine (1/1) adduct and the Xray crystal structure analysis of the resulting dimethylethylammonium bromide, [ $\mathrm{HNEtMe}_{2}$ ]Br. The synthesis of triferrocenylboroxine and of the title compound, (I), was achieved by the hydrolysis of the 1-(dibromoboryl)ferrocenedimethylethylammonium (1/1) adduct, as indicated in the reaction scheme below.


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There is a strong hydrogen bond linking the NH group and the $\mathrm{Br}^{-}$ion. Furthermore, the packing is stabilized by several weak $\mathrm{C}-\mathrm{H} \cdots \mathrm{Br}$ hydrogen bonds.

## Experimental

The title compound, (I), was obtained by stirring a solution of 0.107 g 1-(dibromoboryl)ferrocene-dimethylethylamine (1/1) in 10 ml $\mathrm{CH}_{2} \mathrm{Cl}_{2}$ in the presence of $0.05 \mathrm{ml} \mathrm{H}_{2} \mathrm{O}$ at ambient temperature. The solution was subjected to gas-phase diffusion of diethyl ether, to yield colourless crystals of $\left[\mathrm{HNEtMe}{ }_{2}\right] \mathrm{Br}$. The NMR spectra were recorded on a Bruker AM 250 spectrometer. [ $\mathrm{HNEtMe} \mathrm{e}_{2}$ ]Br: ${ }^{1} \mathrm{H}$ NMR $\left(\mathrm{CDCl}_{3}\right.$, internal TMS $): \delta 1.45\left(t, \mathrm{CH}_{3},{ }^{3} J=7.2 \mathrm{~Hz}\right), 2.78\left(d, 2 \mathrm{NCH}_{3},{ }^{3} J=\right.$ $5.3 \mathrm{~Hz}), 3.14\left(d q, \mathrm{NCH}_{2},{ }^{3} J=7.2 \mathrm{~Hz},{ }^{3} J=5.3 \mathrm{~Hz}\right), 11.26(m, \mathrm{NH})$. ${ }^{13} \mathrm{C}\left\{{ }^{1} \mathrm{H}\right\}$ NMR $\left(\mathrm{CDCl}_{3}\right.$, internal TMS): $\delta 10.0\left(s, \mathrm{CH}_{3}\right), 42.6(s$, $2 \mathrm{NCH}_{3}$ ), $53.3\left(s, \mathrm{NCH}_{2}\right)$.

## Crystal data

$\mathrm{C}_{4} \mathrm{H}_{12} \mathrm{~N}^{+} \cdot \mathrm{Br}^{-}$
$M_{r}=154.06$
Monoclinic, $P 2_{\AA} / c$
$a=7.0797(8) \AA$
$b=7.3352(9) \AA$
$c=14.1594(15) \AA$
$\beta=100.030(9)^{\circ}$
$V=724.07(14) \AA^{3}$
$Z=4$
$D_{x}=1.413 \mathrm{Mg} \mathrm{m}^{-3}$
$M_{r}=154.06$
Monoclinic, $P 2_{1} / c$
$b=7.3352$ (9) $\AA$
$c=14.1594$ (15) $\AA$
$V=724.07(14) \AA^{3}$
$Z=4$

## Data collection

| Stoe IPDS-II two-circle | 1675 independent reflections |
| :--- | :--- |
| $\quad$ diffractometer | 1171 reflections with $I>2 \sigma(I)$ |
| $\omega$ scans | $R_{\text {int }}=0.063$ |
| Absorption correction: multi-scan | $\theta_{\max }=27.8^{\circ}$ |
| $\quad$ (MULABS; Spek, 1990; Blessing, | $h=-9 \rightarrow 9$ |
| 1995) | $k=-8 \rightarrow 9$ |
| $T_{\min }=0.381, T_{\max }=0.485$ | $l=-18 \rightarrow 18$ |
| 5255 measured reflections |  |

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.033$
$w R\left(F^{2}\right)=0.072$
$S=0.85$
1675 reflections
78 parameters

Mo $K \alpha$ radiation
Cell parameters from 1423
reflections
$\theta=3.5-24.2^{\circ}$
$\mu=5.57 \mathrm{~mm}^{-1}$
$T=173$ (2) K
Block, colourless
$0.19 \times 0.17 \times 0.13 \mathrm{~mm}$

1675 independent reflections
1171 reflections with $I>2 \sigma(I)$
$R_{\text {int }}=0.063$
$\theta_{\text {max }}=27.8^{\circ}$
$k=-8 \rightarrow 9$
$l=-18 \rightarrow 18$

H atoms treated by a mixture of independent and constrained refinement
$w=1 /\left[\sigma^{2}\left(F_{o}{ }^{2}\right)+(0.032 P)^{2}\right]$
where $P=\left(F_{o}{ }^{2}+2 F_{c}{ }^{2}\right) / 3$
$(\Delta / \sigma)_{\max }=0.029$
$\Delta \rho_{\text {max }}=0.32 \mathrm{e}^{-3}$
$\Delta \rho_{\text {min }}=-0.56 \mathrm{e}^{\AA^{-3}}$

Table 1
Hydrogen-bonding geometry $\left(\AA^{\circ}{ }^{\circ}\right)$.

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| $\mathrm{~N} 1-\mathrm{H} 1 \cdots \mathrm{Br} 1$ | $0.85(6)$ | $2.37(6)$ | $3.217(3)$ | $173(5)$ |
| $\mathrm{C} 2-\mathrm{H} 2 A \cdots \mathrm{Br}^{1}{ }^{\mathrm{i}}$ | 0.98 | 2.90 | $3.819(5)$ | 156 |
| $\mathrm{C} 1-\mathrm{H} 1 A \cdots \mathrm{Br}^{\mathrm{ii}}$ | 0.98 | 2.88 | $3.849(4)$ | 171 |
| $\mathrm{C} 2-\mathrm{H} 2 B \cdots \mathrm{Br}^{\text {iii }}$ | 0.98 | 3.03 | $3.913(5)$ | 151 |
| $\mathrm{C} 2-\mathrm{H} 2 C \cdots \mathrm{Br}^{\text {iv }}$ | 0.98 | 3.05 | $3.931(5)$ | 150 |
| $\mathrm{C} 1-\mathrm{H} 1 C \cdots \mathrm{Br}^{\mathrm{iii}}$ | 0.98 | 3.00 | $3.895(5)$ | 152 |

Symmetry codes: (i) $-x, y-\frac{1}{2}, \frac{1}{2}-z$; (ii) $1-x, 1-y, 1-z$; (iii) $-x, 1-y, 1-z$; (iv) $x, y-1, z$.

The initial cell determination yielded an orthorhombic $C$-centred cell with $a=18.197 \AA, b=21.698 \AA, c=7.335 \AA$ and $V=2896.1 \AA^{3}$. After data collection, the $R_{\text {int }}$ value for the orthorhombic setting (0.474) indicated that the symmetry should be lowered to monoclinic $P$ with an $R_{\text {int }}$ of 0.063 . Structure solution and refinement was


Figure 1
Perspective view of the title compound with the atom numbering; displacement ellipsoids are drawn at the $50 \%$ probability level. The major component of the disordered ethyl group (C3 and C 4 ) is drawn with full bonds, the minor component ( $\mathrm{C}^{\prime}$ and $\mathrm{C} 4^{\prime}$ ) with open bonds.
successful in $P 2_{1} / c$. The ethyl group is disorded over two sites. The site-occupation factors refined to 0.54 (1):0.46 (1). H atoms bonded to C atoms were refined with fixed individual displacement parameters $\left[U_{\text {iso }}(\mathrm{H})=1.2 U_{\text {eq }}(\mathrm{C})\right.$ or $\left.1.5 U_{\text {eq }}\left(\mathrm{C}_{\text {methyl }}\right)\right]$, using a riding model, with $\mathrm{C}-\mathrm{H}=0.99 \AA$ and methyl $\mathrm{C}-\mathrm{H}=0.98 \AA$. The H atom bonded to the N atom was freely refined.

Data collection: $X$ - $A R E A$ (Stoe \& Cie, 2001); cell refinement: $X$ $A R E A$; data reduction: $X-A R E A$; program(s) used to solve structure: SHELXS97 (Sheldrick, 1990); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: $X P$ in SHELXTL-Plus (Sheldrick, 1991); software used to prepare material for publication: SHELXL97.

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