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Key indicators

Single-crystal X-ray study T = 93 K Mean σ (N–C) = 0.002 Å R factor = 0.010 wR factor = 0.024 Data-to-parameter ratio = 14.4

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

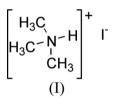
Redetermination of trimethylammonium iodide at 93 K

The crystal structure of the title compound, $C_3H_{10}N^+$ ·I⁻, originally determined by Sheldrick & Sheldrick [*Acta Cryst.* (1970), B**26**, 1334–1338], has been redetermined at 93 K. Both ions lie on a mirror plane. They form N-H···I hydrogenbonded units in the crystal structure.

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Comment

The title compound, (I), was inadvertently formed as a byproduct from a reaction investigating the synthesis of aminoborane derivatives in tetrahydrofuran. The less soluble trimethylammonium iodide crystallized out as colourless crystals, leaving the basic cyclic aminoborane in solution. Although this reaction has not been reported previously, reactions of primary and secondary amines with boron halides have been documented (Nöth *et al.*, 1964).



The compound has been studied before at room temperature by film methods (Sheldrick & Sheldrick, 1970, and references therein). We present here a 'state of the art' lowtemperature data set with a total data collection time of less

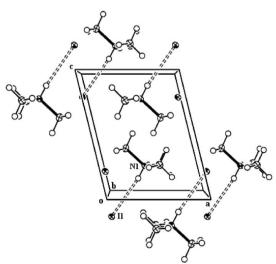


Figure 1

An ORTEP-3 (Farrugia, 1997) view, down the b axis, of the unit-cell contents of (I), shown with 50% probability displacement ellipsoids. $N-H\cdots I$ hydrogen bonds are shown as dashed lines.

© 2006 International Union of Crystallography All rights reserved than 3 h and conventional R factor of 0.010, which is reflected in the ability to refine H atoms in the presence of the much larger scattering I atom given well defined crystals. I, NH and one CH lie on a mirror plane. The structure contains hydrogen-bonded units (Table 2 and Fig. 1), as indicated in the earlier study. The $N-H\cdots I$ angle, expected to be near linear, compares with the reported range for dimethylamino iodide structures (Cambridge Structural Database, Version 5.27, update of May 2006; Allen, 2002) of 157-179° [e.g. 175° in Schneider & Schier (2004)]. The film determination geometry, even without allowance for the different temperature of data collection, is statistically identical although of considerably lower resolution.

Experimental

Iodine (6.61 g, 0.026 mol) was added in small aliquots to a solution of trimethylamine borane (1.9 g, 0.026 mol) containing ethylenediamine (0.78 g, 0.013 mol) in tetrahydrofuran (100 ml). On concentration of the solution, suitable crystals were deposited.

Crystal data

$C_{3}H_{10}N^{+}\cdot I^{-}$	Z = 2
$M_r = 187.02$	$D_x = 1.969 \text{ Mg m}^{-3}$
Monoclinic, $P2_1/m$	Mo $K\alpha$ radiation
a = 5.5309 (7) Å	$\mu = 4.94 \text{ mm}^{-1}$
$b = 8.2737 (10) \text{\AA}$	T = 93 (2) K
c = 7.1000 (9) Å	Block, colourless
$\beta = 103.8010 \ (10)^{\circ}$	$0.37 \times 0.18 \times 0.16$
V = 315.52 (7) Å ³	
Data collection	

Bruker SMART CCD area-detector diffractometer ω and ω scans Absorption correction: multi-scan (Blessing, 1995) $T_{\min} = 0.304, T_{\max} = 0.454$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.010$ $wR(F^2) = 0.024$ S = 1.23735 reflections 51 parameters All H-atom parameters refined 1741 measured reflections 735 independent reflections 730 reflections with $I > 2\sigma(I)$ $R_{\rm int} = 0.011$ $\theta_{\rm max} = 27.9^\circ$

mm

 $w = 1/[\sigma^2(F_0^2) + (0.0077P)^2]$ + 0.1167P] where $P = (F_0^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\rm max} = 0.001$ $\Delta \rho_{\rm max} = 0.34 \text{ e} \text{ Å}^{-3}$ $\Delta \rho_{\rm min} = -0.25 \text{ e} \text{ Å}^{-3}$ Extinction correction: SHELXL97 Extinction coefficient: 0.045 (2)

Table 1

Selected geometric parameters (Å, °).

N1-C1	1.498 (3)	N1-H3	0.81 (3)
N1-C2	1.4977 (18)	C1-H1	0.94 (3)
C1-N1-C2 C1-N1-H3	111.26 (11) 105 (2)	C2-N1-H3	108.7 (11)

Table 2

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdots A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$N1 - H3 \cdots I1^i$	0.81 (3)	2.70 (3)	3.500 (2)	173 (3)
Symmetry code: (i) $x + 1, y, z + 1$.			

All H atoms were refined with isotropic displacement parameters [N-H = 0.81 (3) Å and C-H = 0.94 (2)-1.00 (2) Å].

Data collection: SMART (Siemens, 2001); cell refinement: SAINT (Siemens, 2001); data reduction: SAINT and SADABS (Sheldrick, 1996); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEP-3 (Farrugia, 1997); software used to prepare material for publication: SHELXL97.

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