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Comment

It is known that 2-aminoperimidylammonium chloride is an excellent reagent for the removal of sulfate ions (Stephen, 1970). However, the synthesis of this compound is inconvenient, and the purification procedure described by Stephen (1970) does not remove every trace of sulfur-containing impurities from the product (McClure, 1973). A synthetic method for the preparation of the bromide salt, free of sulfate impurities, is given by McClure (1973) and is suggested to be a simpler preparation of the sulfate-removing reagent. For experimental purposes, it was desired by one of the authors (CEM) to prepare a polymer of this compound to remove sulfate from water. It was also desired that the polymer be easy to handle and reusable. Such a polymer was prepared, but it retained sulfate too strongly to be regenerated efficiently. Subsequent work on the polymer has provided a way to regenerate the polymer and will be reported at a future date.

The crystal structure of the title compound, (I), was determined by three-dimensional X-ray diffraction analysis in order to study possible sulfate binding sites. An unsuccessful attempt was made to complex the title compound with sulfate using a gel-permeation technique in order to determine exactly how and where the compound binds to sulfate.



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2-Aminoperimidinium Bromide Dihydrate

STACY M. BAKER,^{*a*} RUSSELL G. BAUGHMAN,^{*a**} CLIFTON E. MELOAN^{*b*} AND PETER MUMBA^{*b*}

^aDivision of Science, Northeast Missouri State University, Kirksville, MO 63501-0828, USA, and ^bDepartment of Chemistry, Kansas State University, Manhattan, KS 66506, USA. E-mail: sc05@nemomus.edu

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Abstract

The title compound, $C_{11}H_{10}N_3^+$.Br⁻.2H₂O, is essentially planar along the aminoperimidine skeleton, with an r.m.s. deviation of 0.009 Å and a maximum deviation of 0.017 Å. Due to the presence of two water molecules, extensive hydrogen bonding exists in the crystal. Hydrogen bonds *via* the two water molecules join the Br⁻ ion to both the amine N atom and one of the ring N atoms. The other ring N atom and the amine N atom also form hydrogen bonds to the Br⁻ ion from the second molecule in the unit cell. The plane defined by all the non-H atoms in the aminoperimidine skeleton displayed a maximum deviation of 0.017 Å for atom N1 and an r.m.s. deviation of



Fig. 1. View of the title compound. Displacement ellipsoids are shown at 50% probability levels and H atoms are drawn as small spheres of arbitrary radii.

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Br

01

O2 N1

N2

N3 CI C2 C3 C4 Ċ5

C6 C7

C8 C9

C10

C11

0.009 Å. Thus, the cation possesses near $C_{2\nu}$ symmetry. Other deviations were Br at 1.939(4), O1 at -0.124(4)and O2 at 0.579 (4) Å.

Due to the presence of the two water molecules, several hydrogen bonds occur. As shown in Fig. 1, these hydrogen bonds join the Br⁻ ion to the N2 and N3 atoms via the two water molecules. The H atoms bound to the N1 and N3 atoms also bind to the Br⁻ ion translated one unit in the y direction.

Experimental

The title compound was obtained from the Aldrich Chemical Co. and crystals were grown by slow evaporation of an aqueous solution.

Crystal data

$C_{11}H_{10}N_3^+.Br^2H_2O$	Mo $K\alpha$ radiation		lessed accur	atuia navanatana	(Å ?)
$M_r = 300.16$	$\lambda = 0./10/3 \text{ A}$	Table 2. Set	ieciea geom	erric parameters	(A,)
Triclinic	Cell parameters from 25	NI-CI	1.339 (4)	N1-C10	1.411 (4)
P1	reflections	N2C1	1.323 (4)	N2C2	1.404 (4)
a = 7.1560 (10) Å	$\theta = 5.6 - 12.2^{\circ}$	N3—C1	1.320 (4)	C2—C3	1.369 (4)
h = 8599(4)Å	$\mu = 3.208 \text{ mm}^{-1}$	C2C11	1.414 (4)	C3-C4	1.399 (5)
c = 11.060(4) Å	T = 288(2) K	C4—C5	1.368 (5)	C5C6	1.413 (5)
t = 11.000 (4) R	I = 200 (2) I R	C6C11	1.414(4)	C^{0}	1.413 (3)
$\alpha = 100.02(3)$		C_{1}	1.362 (3)	C_{1}	1.407(5) 1.408(4)
$\beta = 104.81(2)^{\circ}$	$0.58 \times 0.28 \times 0.19$ mm	C9-C10	1.505 (4)		1.400 (4)
$\gamma = 96.20(3)^{\circ}$	Colorless	N3—C1—N2	121.6 (3)	C5-C6-C11	118.1(3)
$V = 639.6 (4) \text{ Å}^3$		N2 - C1 - N1	120.0 (3)	$C_1 = C_0 = C_1$	118.0(3)
Z = 2		$C_3 = C_2 = C_1$	120.7(3)	$C_{1} = C_{1} = C_{1}$	121.4(3) 121.4(3)
$D_{\rm c} = 1.559 {\rm Mg}{\rm m}^{-3}$		$C_2 = C_3 = C_4$	120 1 (3)	C_{11} C_{10} N_{11}	121.4(3) 116.8(3)
$D_x = 1.557$ Mg m		$C_{4} = C_{5} = C_{0}$	123.9 (3)	C10-C11-C6	119.7 (3)
D_m not measured		C8C7C6	120.7 (3)	C1 - N1 - C10	122.9 (3)
		C10-C9-C8	118.8 (3)	CI-NI-HIA	118.6 (2)
		C9-C10-N1	121.9 (3)	C10—N1—H1A	118.6(2)
Data collection		C10-C11-C2	120.2 (3)	C1-N2-C2	123.0 (3)
Siemens P3 diffractometer	$R_{\rm int} = 0.0307$	C2-C11-C6	120.1 (3)	C1-N2-H2A	118.6 (2)
	$A = 25.05^{\circ}$	N3-C1-N1	118.5 (3)	C2—N2—H2A	118.5 (2)
0120 scans	$v_{\text{max}} = 25.05$	C3—C2—N2	122.1 (3)	C1—N3—H3A	118.9 (2)
Absorption correction:	$h = -8 \rightarrow 8$	N2-C2-C11	117.2 (3)	C1 - N3 - H3B	121.1(2)
six ψ scans, 10° steps	$k = -10 \rightarrow 10$	C5-C4-C3	122.2 (3)	H3A—N3—H3B	120.0
(Siemens, 1991a)	$l = -13 \rightarrow 13$	Hydrogen-bonding geometry			
$T_{\rm min} = 0.16, \ T_{\rm max} = 0.54$	3 standard reflections	Br H11	2 348 (7)	Br. H3A'	2 627 (3)
4501 measured reflections	monitored every 50	Br H13	2 358 (3)	$01 \cdots H2A$	1.854 (3)
2266 independent reflections	reflections	Br. · ·HIA'	2.491 (3)	$O2 \cdot \cdot \cdot H3B$	2.153 (4)
1720 observed reflections	intensity decay: average	U11 D- U12	64(1)	01H24N2	1747(1)
	2.5% in $-(D)$	$H1A^1 \dots Br \dots H3A^1$	51 32 (8)	$N3 - H34 \cdots Br^{H}$	153 31 (6)
$[I > 2\sigma(I)]$	$2.5\% \prod O(1)$	Br. Ol. H2A	128.0 (2)	$O2 \cdot \cdot H3B - N3$	149.5 (1)
		$H3B \cdots O2 - H13$	113 (2)	Br···H11—O1	168 (2)
		NI-HIA···Br"	159.38 (5)	Br···H13-O2	178 (3)
D (

Symmetry codes: (i) x, y - 1, z; (ii) x, 1 + y, z.

Re	finemen
nc.	memen

Refinement on F^2 R(F) = 0.0364 $wR(F^2) = 0.0777$ S = 1.0362257 reflections 166 parameters H atoms riding (included in least-squares refinement with C-H 0.96, O-H 1.00 and N-H 0.90 Å)

 $w = 1/[\sigma^2(F_o^2) + (0.0416P)^2$ + 0.1304P] where $P = (F_0^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\rm max} < 0.001$ $\Delta \rho_{\rm max} = 0.289 \ {\rm e} \ {\rm \AA}^{-3}$ $\Delta \rho_{\rm min} = -0.280 \ {\rm e} \ {\rm \AA}^{-3}$ Atomic scattering factors from International Tables for Crystallography (1992, Vol. C, Tables 4.2.6.8 and 6.1.1.4)

Data collection: P3/P4-PC Diffractometer Program (Siemens, 1991a). Cell refinement: P3/P4-PC Diffractometer Program. Data reduction: XDISK (Siemens, 1991b). Program(s) used to solve structure: SHELXS86 (Sheldrick, 1990a). Program(s) used to refine structure: SHELXL93 (Sheldrick, 1993). Molecular graphics: SHELXTL/PC (Sheldrick, 1990b). Software used to prepare material for publication: SHELXTL/PC and SHELXL93.

Table 1. Fractional atomic coordinates and equivalent isotropic displacement parameters $(Å^2)$

$U_{eq} =$	(1/3)	$\Sigma_i \Sigma_j \iota$	l _{ij} a*a	$\mathbf{a}_i \cdot \mathbf{a}_j$.
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• • • • • • • • • • • • • • • • • • • •					
x	y	z	U_{eq}		
0.3419(1)	-0.2366 (1)	0.5796(1)	0.060 (1)		
0.1042 (4)	0.0100 (3)	0.7289(3)	0.061(1)		
0.2427 (4)	0.0931 (3)	().4680(3)	0.062(1)		
0.2870(4)	0.5927 (3)	0.8210(2)	().039(1)		
0.2092 (4)	0.3313 (3)	0.8368 (2)	0.040(1)		
0.2334 (4)	0.3913 (3)	0.6455 (3)	0.053(1)		
0.2424 (5)	0.4364 (4)	0.7675 (3)	0.038(1)		
0.2161 (4)	0.3754 (4)	0.9664 (3)	0.037(1)		
0.1798 (5)	0.2649 (4)	1.0363 (3)	0.046 (1)		
0.1894 (5)	0.3180(5)	1.1651 (4)	0.053(1)		
0.2329 (5)	0.4767 (5)	1.2231 (3)	0.051(1)		
0.2704 (4)	0.5932 (4)	1.1530 (3)	0.040(1)		
0.3151 (5)	0.7597 (4)	1.2050 (3)	0.050(1)		
0.3466 (5)	0.8655 (4)	1.1311 (3)	0.052(1)		
0.3371 (5)	0.8135 (4)	1.0015 (3)	0.045(1)		
().2970 (4)	0.6533 (4)	().9499 (3)	0.037(1)		
().2623(4)	0.5405 (3)	1.0231 (3)	0.033(1)		

Lists of structure factors, anisotropic displacement parameters, Hatom coordinates and complete geometry have been deposited with the IUCr (Reference: CR1182). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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1-(4-Nitrophenyl)-4-piperidinol

DAVID W. TOMLIN,^{*a*} TIMOTHY J. BUNNING,^{*b*} GARY E. PRICE,^{*c*} ALBERT V. FRATINI^{*d*} AND W. WADE ADAMS^{e*}

^aTechnical Management Concepts, Inc, PO Box 340345, Beavercreek, OH 45434, USA, ^bScience Applications International Corporation, 101 Woodman Drive, Suite 103, Dayton, OH 45431, USA, ^cResearch Institute, University of Dayton, Dayton, OH 45469, USA, ^dDepartment of Chemistry, University of Dayton, Dayton, OH 45469, USA, and ^eWright Laboratory, Materials Directorate, Wright-Patterson AFB, OH 45433, USA. E-mail: adamsww@ml.wpafb.af.mil

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Abstract

The title compound, $C_{11}H_{14}N_2O_3$, is a non-linear optical chromophore. The piperidinol ring is in a chair conformation. The C—N—C fragment of the piperidinol moiety is nearly coplanar with the nitrophenyl ring system. The molecular stacking allows hydrogen bonding between the piperidinol hydroxy group and the nitro group.

Comment

The present study is part of a program designed to investigate the structure-property relationships among nonlinear optical (NLO) materials. Organic NLO materials have potential for use in optical limiting devices for applications in sensor protection, optical storage devices and waveguides (Prasad & Williams, 1991). Owing to their versatility of chemical synthesis, organic materials are increasingly being investigated for their NLO properties (Prasad & Williams, 1991; Hann & Bloor, 1991; Chemla & Zyss, 1987; Miyata, 1992). Organic structures are versatile because they may be grown as single crystals, as Langmuir–Blodgett films or incorporated as chromophoric mesogens in polymer systems (Prasad & Williams, 1991).

Second-order NLO effects are highly dependent on the geometrical arrangement of the molecules in the crystal phase. For a crystalline material to exhibit second-order optical non-linearity, the crystal must be non-centrosymmetric (Prasad & Williams, 1991; Hann & Bloor, 1989). The present crystals, (I), are centrosymmetric and therefore do not exhibit NLO behavior. The production of a non-centrosymmetric material by using the compound as a chromophoric mesogen on a polymer backbone is being investigated. Molecular substitution on the piperidinol moiety is also being investigated, but the substituent must not interfere with the molecular features responsible for the optical non-linearity (Zyss & Oudar, 1982).



Bond lengths and angles in the nitrophenyl group are consistent with those of both nitrobenzene itself (Trotter, 1959) and the nitrophenyl moiety found in related compounds (Fkyerat *et al.*, 1995; Ishihara *et al.*, 1992; Hall *et al.*, 1986; Twieg & Dirk, 1986). Bond lengths and angles in the piperidinol ring are consistent with those of piperidine (Suwinska, Palenik & Gerdil, 1986). The piperidinol moiety adopts a chair conformation



Fig. 1. The molecular structure of the title compound showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level.

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