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Key indicators

Single-crystal X-ray study
 $T = 120$ K
Mean $\sigma(\text{C}-\text{C}) = 0.003$ Å
 R factor = 0.022
 wR factor = 0.055
Data-to-parameter ratio = 19.4For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.Propane-1,3-diaminium hydrogenarsenate
monohydrate

The title compound, $(\text{C}_3\text{H}_{12}\text{N}_2)[\text{HAsO}_4]\cdot\text{H}_2\text{O}$, contains a network of propane-1,3-diaminium cations, hydrogenarsenate anions [mean $\text{As}-\text{O} = 1.687$ (2) Å] and water molecules. The crystal packing involves anion-to-anion and water-to-anion $\text{O}-\text{H}\cdots\text{O}$ hydrogen bonds, resulting in infinite chains containing the unusual $R_3^3(10)$ graph-set motif. Cation-to-anion and cation-to-water $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds generate a three-dimensional overall structure.

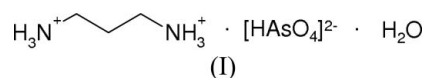
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Comment

The title compound, $(\text{C}_3\text{H}_{12}\text{N}_2)[\text{HAsO}_4]\cdot\text{H}_2\text{O}$, (I) (Fig. 1), was prepared as part of our ongoing structural studies of hydrogen-bonding interactions in protonated-amine (di)hydrogen arsenates (Lee & Harrison, 2003*a*; Wilkinson & Harrison, 2004; Todd & Harrison, 2005). In particular, (I) complements propane-1,3-diaminium bis(dihydrogenarsenate), $(\text{C}_3\text{H}_{12}\text{N}_2)[\text{H}_2\text{AsO}_4]_2$ (Wilkinson & Harrison, 2005), prepared under different pH conditions.



The $[\text{HAsO}_4]^{2-}$ hydrogenarsenate group in (I) has normal tetrahedral geometry [mean $\text{As}-\text{O} = 1.687$ (2) Å], with the protonated $\text{As1}-\text{O4}$ vertex showing its usual lengthening relative to the unprotonated $\text{As}-\text{O}$ bonds (Table 1). The propane-1,3-diaminium cation shows no unusual geometrical features.

As well as electrostatic attractions, the component species in (I) interact by means of a network of $\text{O}-\text{H}\cdots\text{O}$ and $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds (Table 2). The $[\text{HAsO}_4]^{2-}$ units and water molecules are linked into polymeric chains (Fig. 2) propagating along [010] by way of anion-to-anion $\text{O4}-\text{H1}\cdots\text{O2}^i$ and water-to-anion $\text{O5}-\text{H14}\cdots\text{O1}$ and $\text{O5}-$

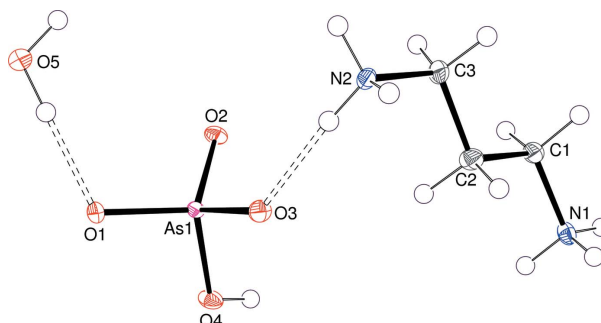


Figure 1

A view of (I) with the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level. Hydrogen bonds are indicated by dashed lines.

H15...O2ⁱⁱ bonds (Table 2). This arrangement results in an unusual $R_3^3(10)$ graph-set (Bernstein *et al.*, 1995) motif. The As1...As1ⁱ separation is 4.7991 (3) Å.

The organic species interacts with the hydrogenarsenate/water chains by way of six N—H...O hydrogen bonds [mean H...O = 1.89 Å, mean N—H...O = 171° and mean N...O = 2.793 (2) Å]. One of the acceptor O atoms is part of a water molecule, and the other five are parts of hydrogenarsenate groups. This hydrogen-bonding scheme results in a three-dimensional network (Fig. 3).

The hydrogen-bonded hydrogenarsenate/water chains in (I) are different from the motifs seen in related structures. In bis(cycloheptylaminium) hydrogenarsenate monohydrate (Todd & Harrison, 2005) and bis(benzylammonium) hydrogenarsenate monohydrate (Lee & Harrison, 2003c), hydrogen-bonded dimers of [HAsO₄]²⁻ units occur, with the dimers bridged into double chains by intervening water molecules. In the unhydrated piperidinium dihydrogenarsenate (Lee & Harrison, 2003b) and *t*-butylammonium dihydrogenarsenate (Wilkinson & Harrison, 2004), single chains of [H₂AsO₄]⁻ anions occur with each adjacent dihydrogenarsenate pair linked by a pair of hydrogen bonds. In propane-1,3-diaminium bis(dihydrogenarsenate) (Wilkinson & Harrison, 2005), the same organic cation as found in (I) is combined with dihydrogenarsenate [H₂AsO₄]⁻ groups, with the latter forming double chains.

Experimental

0.5 M aqueous propane-1,3-diamine solution (10 ml) was added to 0.5 M aqueous H₃AsO₄ solution (10 ml) to result in a clear solution. Aqueous ammonia was added to this solution to raise the pH to about 12, which is beyond the second end-point for H₃AsO₄ (*i.e.* the predominant species is [HAsO₄]²⁻). Platy crystals of (I) grew as the water evaporated over the course of a few days.

Crystal data

(C ₃ H ₁₂ N ₂)[HAsO ₄]·H ₂ O	$D_x = 1.778 \text{ Mg m}^{-3}$
$M_r = 234.09$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/c$	Cell parameters from 2043 reflections
$a = 7.1327 (2) \text{ \AA}$	$\theta = 2.9\text{--}27.5^\circ$
$b = 16.8046 (6) \text{ \AA}$	$\mu = 3.87 \text{ mm}^{-1}$
$c = 7.9402 (2) \text{ \AA}$	$T = 120 (2) \text{ K}$
$\beta = 113.253 (2)^\circ$	Plate, colourless
$V = 874.42 (5) \text{ \AA}^3$	$0.32 \times 0.24 \times 0.03 \text{ mm}$
$Z = 4$	

Data collection

Nonius KappaCCD diffractometer	1804 reflections with $I > 2\sigma(I)$
ω and φ scans	$R_{\text{int}} = 0.036$
Absorption correction: multi-scan (SADABS; Bruker, 1999)	$\theta_{\text{max}} = 27.5^\circ$
$T_{\text{min}} = 0.370, T_{\text{max}} = 0.892$	$h = -9 \rightarrow 8$
11562 measured reflections	$k = -20 \rightarrow 21$
2002 independent reflections	$l = -10 \rightarrow 10$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0218P)^2 + 0.7899P]$
$R[F^2 > 2\sigma(F^2)] = 0.022$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.055$	$(\Delta/\sigma)_{\text{max}} = 0.001$
$S = 1.05$	$\Delta\rho_{\text{max}} = 0.49 \text{ e \AA}^{-3}$
2002 reflections	$\Delta\rho_{\text{min}} = -0.53 \text{ e \AA}^{-3}$
103 parameters	Extinction correction: SHELXL97
H-atom parameters constrained	Extinction coefficient: 0.0032 (6)

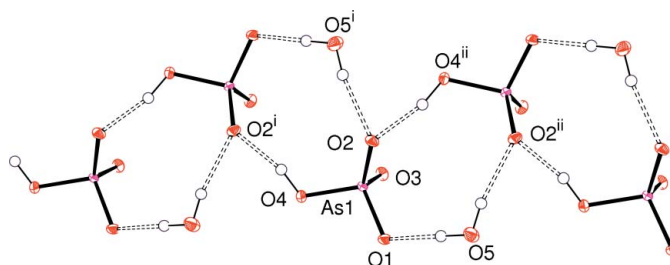


Figure 2 Detail of a hydrogen-bonded (dashed lines) hydrogenarsenate/water chain in (I).

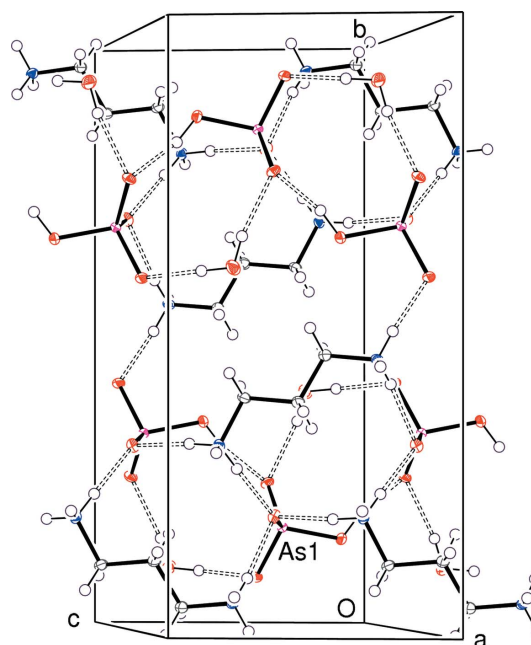


Figure 3 The crystal packing of (I). Dashed lines indicate hydrogen bonds.

Table 1 Selected geometric parameters (Å, °).

As1—O1	1.6612 (14)	As1—O3	1.6814 (14)
As1—O2	1.6746 (13)	As1—O4	1.7302 (13)
N1—C1—C2—C3	175.33 (16)	C1—C2—C3—N2	175.49 (16)

Table 2 Hydrogen-bond geometry (Å, °).

D—H...A	D—H	H...A	D...A	D—H...A
O4—H1...O2 ⁱ	0.93	1.71	2.6207 (19)	166
O5—H14...O1	0.92	1.79	2.709 (2)	177
O5—H15...O2 ⁱⁱ	0.89	1.98	2.858 (2)	169
N1—H2...O1 ⁱⁱⁱ	0.91	1.81	2.711 (2)	173
N1—H3...O3 ⁱ	0.91	1.96	2.855 (2)	166
N1—H4...O5 ^{iv}	0.91	1.90	2.798 (2)	168
N2—H11...O3 ⁱⁱ	0.91	1.90	2.802 (2)	170
N2—H12...O2 ^v	0.91	1.95	2.851 (2)	172
N2—H13...O3	0.91	1.84	2.743 (2)	175

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

The O-bound H atoms were found in difference maps and refined as riding on their carrier O atoms in their as-found relative positions. H atoms bonded to C and N atoms were placed in idealized positions (C–H = 0.99 Å and N–H = 0.91 Å) and refined as riding, allowing for free rotation of the –NH₃ groups. The constraint $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{carrier})$ was applied in all cases.

Data collection: *COLLECT* (Nonius, 1998); cell refinement: *SCALEPACK* (Otwinowski & Minor, 1997); data reduction: *SCALEPACK*, *DENZO* (Otwinowski & Minor, 1997) and *SORTAV* (Blessing, 1995); 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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