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4-Acetylpyridinium hydrogen sulfate

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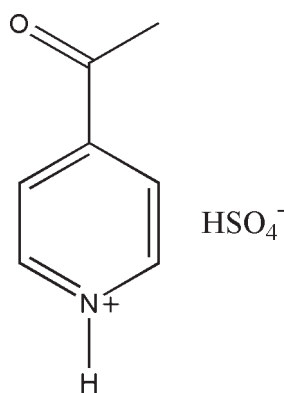
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 Key indicators: single-crystal X-ray study; $T = 298$ K; mean $\sigma(\text{C}-\text{C}) = 0.006$ Å; R factor = 0.053; wR factor = 0.170; data-to-parameter ratio = 16.7.

The crystal structure of the title compound, $\text{C}_7\text{H}_8\text{NO}^+\cdot\text{HSO}_4^-$, consists of $\text{O}-\text{H}\cdots\text{O}$ hydrogen-bonded extended chains of hydrogen sulfate anions. Each hydrogen sulfate anion is furthermore connected to one 4-acetylpyridinium cation *via* a hydrogen bond of the $\text{N}-\text{H}\cdots\text{O}$ type.

Related literature

For the synthesis of 4-acetylpyridine, see: Piner *et al.* (1934). For the crystal structure of an adduct of 4-acetylpyridine with pentachlorophenol, see: Majerz *et al.* (1991). For the crystal structures of Zn and Ni complexes of 4-acetylpyridine, see: Pang *et al.* (1994); Steffen *et al.* (1977).



Experimental

Crystal data

 $\text{C}_7\text{H}_8\text{NO}^+\cdot\text{HSO}_4^-$
 $M_r = 219.21$

 Orthorhombic, $P2_12_12_1$
 $a = 4.6454$ (9) Å

 $b = 9.597$ (2) Å

 $c = 21.310$ (4) Å

 $V = 950.1$ (3) Å³
 $Z = 4$

 Mo $K\alpha$ radiation

 $\mu = 0.34$ mm⁻¹
 $T = 298$ K

 $0.20 \times 0.20 \times 0.20$ mm

Data collection

Rigaku SCXmini diffractometer

Absorption correction: multi-scan

 (*CrystalClear*; Rigaku, 2005)

 $T_{\text{min}} = 0.935$, $T_{\text{max}} = 0.935$

9843 measured reflections

2156 independent reflections

 1622 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.077$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.053$
 $wR(F^2) = 0.170$
 $S = 0.93$

2156 reflections

129 parameters

H-atom parameters constrained

 $\Delta\rho_{\text{max}} = 0.38$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.19$ e Å⁻³

Absolute structure: Flack (1983),

860 Friedel pairs

Flack parameter: 0.2 (2)

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{N1}-\text{H1B}\cdots\text{O1}^{\text{i}}$	0.86	1.92	2.772 (5)	172
$\text{O3}-\text{H3}\cdots\text{O2}^{\text{ii}}$	0.82	1.76	2.565 (5)	166

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

Data collection: *CrystalClear* (Rigaku, 2005); cell refinement: *CrystalClear*; data reduction: *CrystalClear*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *PRPKAPPA* (Ferguson, 1999).

The authors are grateful to the starter fund of Southeast University for financial support to buy the X-ray diffractometer.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: IM2126).

References

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supporting information

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4-Acetylpyridinium hydrogen sulfate

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S1. Comment

4-Acetylpyridine may be used as a ligand in coordination compounds *e.g.* with Zn (Steffen & Palenik, 1977) or Ni (Pang *et al.*, 1994). The crystal structure of 4-acetylpyridine together with pentachlorophenol is also known (Majerz *et al.*, 1991).

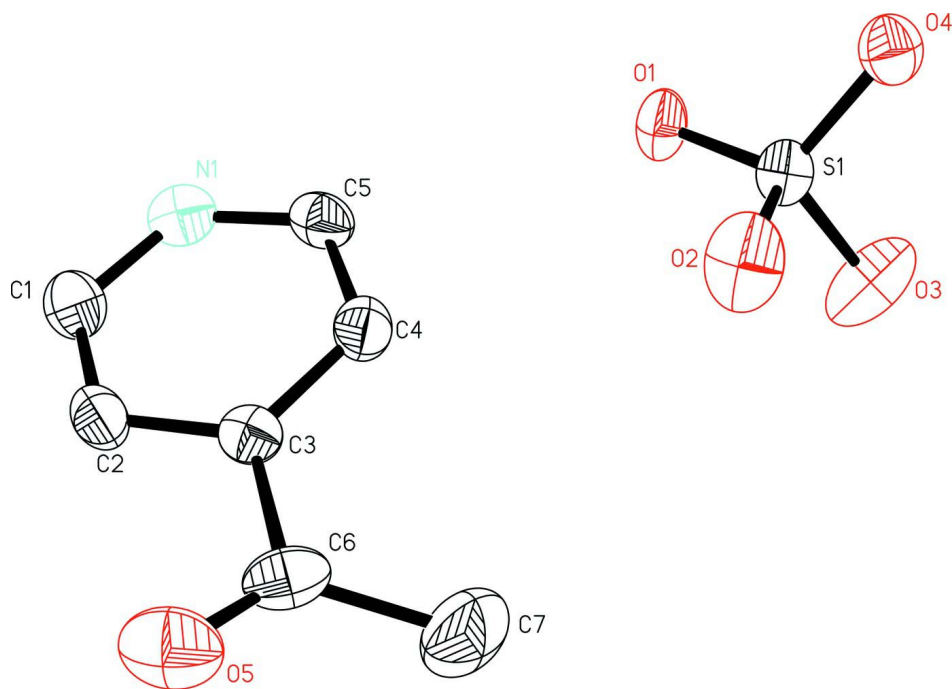
The asymmetric unit of the title compound contains one 4-acetylpyridinium cation and one hydrogen sulfate anion (Fig 1). In the anion, the bond length of S1—O3 is 1.553 (6) Å compared to the average bond length of 1.438 (5) Å of the other S1—O bonds. It is therefore reasonable that the hydrogen atom of hydrogen sulfate is bonded to O3. The supramolecular structure consists of infinite chains of anions with one cation linked to each anion *via* an additional hydrogen bond (N1—H1B···O1 2.774 (8) Å, Fig 2).

S2. Experimental

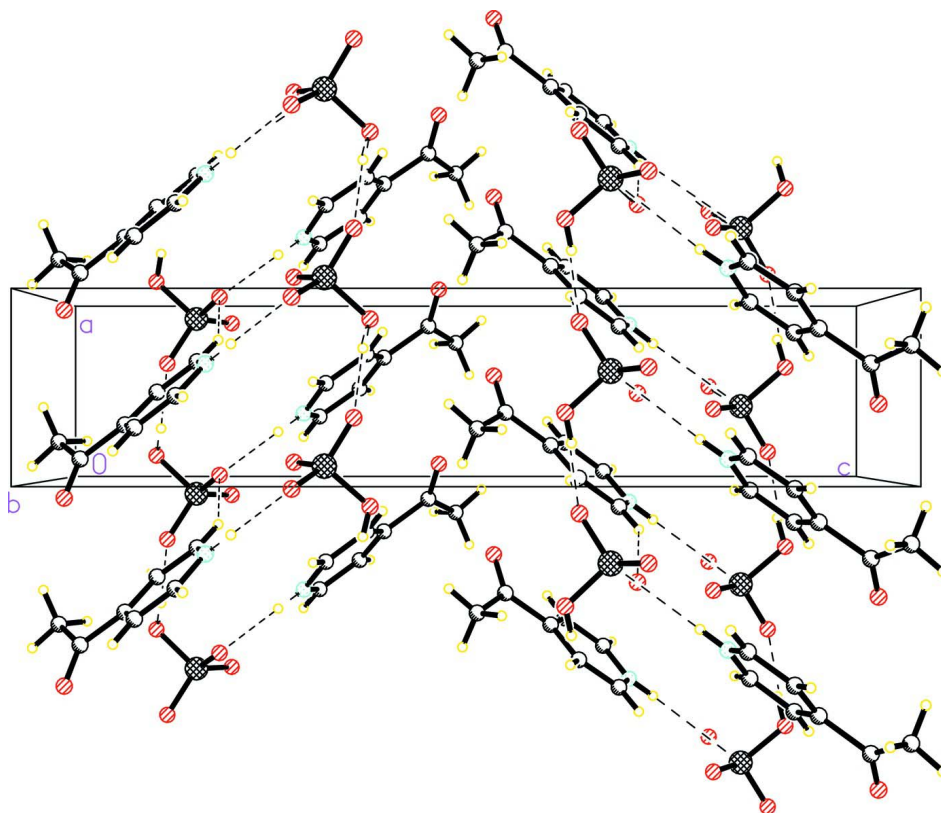
4-Acetylpyridine was obtained according to the method described by Piner (1934). Reaction of equimolar amounts of 4-acetylpyridine and H₂SO₄ produced a precipitate. This was filtered off, dried and dissolved in 96% ethanol from which single crystals were grown by slow evaporation of the solvent at room temperature.

S3. Refinement

The H atom connected to O3 was discernible from difference electron-density map. Nevertheless, it was placed to the ideal position, with S1—O3—H angle tetrahedral, allowing the H atom to ride on the immediately preceding atom O3 and rotate about the S1—O3 bond, refined in a riding atom approximation with a constrained bond length of O—H = 0.82 Å. Positional parameters of the other H atoms were calculated geometrically with C_{ar}—H = 0.93 Å and C_{Me}—H = 0.96 Å and were allowed to ride on the corresponding C atoms with $U_{\text{iso}}(\text{H}) = 1.2 U_{\text{eq}}(\text{C})$. In the absence of significant anomalous dispersion effects, 860 Friedel pairs were merged.

**Figure 1**

Molecular structure of the title compound with the atomic numbering scheme. Displacement ellipsoids are drawn at the 30% probability level and all H atoms have been omitted for clarity.

**Figure 2**

Crystal packing of the title compound, stacking along the *b* axis. Dashed lines indicate hydrogen bonds.

4-Acetylpyridinium hydrogen sulfate

Crystal data

$C_7H_8NO^+ \cdot HSO_4^-$

$M_r = 219.21$

Orthorhombic, $P2_12_12_1$

Hall symbol: P 2ac 2ab

$a = 4.6454(9) \text{ \AA}$

$b = 9.597(2) \text{ \AA}$

$c = 21.310(4) \text{ \AA}$

$V = 950.1(3) \text{ \AA}^3$

$Z = 4$

$F(000) = 456$

$D_x = 1.533 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$

Cell parameters from 4231 reflections

$\theta = 3.6\text{--}27.6^\circ$

$\mu = 0.34 \text{ mm}^{-1}$

$T = 298 \text{ K}$

Prism, colourless

$0.20 \times 0.20 \times 0.20 \text{ mm}$

Data collection

Rigaku SCXmini
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

Detector resolution: $13.6612 \text{ pixels mm}^{-1}$

ω scans

Absorption correction: multi-scan

(*CrystalClear*; Rigaku, 2005)

$T_{\min} = 0.935$, $T_{\max} = 0.935$

9843 measured reflections

2156 independent reflections

1622 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.077$

$\theta_{\max} = 27.5^\circ$, $\theta_{\min} = 3.6^\circ$

$h = -6 \rightarrow 6$

$k = -12 \rightarrow 12$

$l = -27 \rightarrow 27$

*Refinement*Refinement on F^2

Least-squares matrix: full

 $R[F^2 > 2\sigma(F^2)] = 0.053$ $wR(F^2) = 0.170$ $S = 0.93$

2156 reflections

129 parameters

0 restraints

Primary atom site location: structure-invariant
direct methodsSecondary atom site location: difference Fourier
mapHydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained

 $w = 1/[\sigma^2(F_o^2) + (0.1249P)^2 + 1.8027P]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\max} < 0.001$ $\Delta\rho_{\max} = 0.38 \text{ e } \text{\AA}^{-3}$ $\Delta\rho_{\min} = -0.19 \text{ e } \text{\AA}^{-3}$ Absolute structure: Flack (1983), 860 Friedel
pairs

Absolute structure parameter: 0.2 (2)

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R -factors(gt) *etc.* and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
S1	0.9101 (2)	0.18577 (10)	0.16383 (5)	0.0475 (3)
O5	-0.0805 (8)	0.6665 (5)	0.03394 (17)	0.0853 (12)
O1	1.0236 (7)	0.3041 (3)	0.19787 (16)	0.0655 (9)
N1	0.6303 (8)	0.7352 (4)	0.20080 (17)	0.0573 (9)
H1B	0.7336	0.7648	0.2315	0.069*
C3	0.3084 (8)	0.6431 (4)	0.10302 (18)	0.0457 (9)
C1	0.4610 (10)	0.8250 (5)	0.1713 (2)	0.0614 (11)
H1A	0.4549	0.9175	0.1842	0.074*
C6	0.1250 (13)	0.5981 (6)	0.0476 (2)	0.0634 (13)
C5	0.6477 (10)	0.6005 (5)	0.1849 (2)	0.0548 (11)
H5A	0.7674	0.5403	0.2070	0.066*
C4	0.4852 (9)	0.5528 (5)	0.1353 (2)	0.0544 (11)
H4A	0.4949	0.4596	0.1236	0.065*
C2	0.2960 (10)	0.7819 (4)	0.1223 (2)	0.0546 (10)
H2A	0.1759	0.8446	0.1017	0.066*
O2	0.6532 (7)	0.2155 (4)	0.1291 (2)	0.0858 (12)
O3	1.1265 (8)	0.1454 (6)	0.1117 (2)	0.0968 (15)
H3	1.2838	0.1802	0.1194	0.145*
O4	0.8942 (13)	0.0646 (4)	0.2026 (2)	0.1068 (15)
C7	0.2139 (18)	0.4694 (7)	0.0132 (3)	0.106 (2)
H7A	0.0795	0.4510	-0.0200	0.160*
H7B	0.2166	0.3919	0.0417	0.160*
H7C	0.4026	0.4824	-0.0042	0.160*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.0356 (4)	0.0481 (5)	0.0588 (5)	-0.0017 (4)	-0.0011 (4)	-0.0066 (5)
O5	0.058 (2)	0.121 (3)	0.077 (2)	-0.004 (3)	-0.0169 (19)	0.008 (2)
O1	0.069 (2)	0.0471 (16)	0.080 (2)	0.0051 (15)	-0.0109 (17)	-0.0183 (15)
N1	0.046 (2)	0.073 (2)	0.0531 (19)	-0.0074 (18)	0.0000 (17)	-0.0004 (17)
C3	0.0372 (18)	0.054 (2)	0.0459 (19)	-0.0021 (16)	0.0076 (16)	0.0062 (17)
C1	0.063 (3)	0.049 (2)	0.072 (3)	-0.003 (2)	0.002 (2)	0.000 (2)
C6	0.059 (3)	0.083 (3)	0.048 (2)	-0.016 (3)	0.009 (2)	0.003 (2)
C5	0.044 (2)	0.066 (3)	0.054 (2)	0.007 (2)	-0.0010 (19)	0.0119 (19)
C4	0.060 (3)	0.045 (2)	0.058 (2)	0.0030 (19)	0.014 (2)	0.0029 (18)
C2	0.055 (2)	0.047 (2)	0.062 (2)	0.0078 (19)	0.001 (2)	0.0075 (19)
O2	0.0428 (18)	0.096 (3)	0.118 (3)	0.0099 (17)	-0.021 (2)	-0.021 (2)
O3	0.0471 (19)	0.145 (4)	0.098 (3)	-0.010 (2)	0.001 (2)	-0.060 (3)
O4	0.137 (4)	0.073 (2)	0.110 (3)	-0.039 (3)	-0.028 (3)	0.022 (2)
C7	0.128 (6)	0.110 (5)	0.081 (4)	0.002 (5)	-0.020 (4)	-0.039 (4)

Geometric parameters (\AA , $^\circ$)

S1—O4	1.429 (4)	C1—C2	1.360 (6)
S1—O2	1.433 (4)	C1—H1A	0.9300
S1—O1	1.447 (3)	C6—C7	1.495 (8)
S1—O3	1.547 (4)	C5—C4	1.378 (6)
O5—C6	1.194 (7)	C5—H5A	0.9300
N1—C1	1.325 (6)	C4—H4A	0.9300
N1—C5	1.339 (6)	C2—H2A	0.9300
N1—H1B	0.8600	O3—H3	0.8200
C3—C4	1.378 (6)	C7—H7A	0.9600
C3—C2	1.395 (6)	C7—H7B	0.9600
C3—C6	1.519 (6)	C7—H7C	0.9600
O4—S1—O2	114.7 (3)	C7—C6—C3	117.5 (5)
O4—S1—O1	111.6 (2)	N1—C5—C4	118.8 (4)
O2—S1—O1	113.9 (2)	N1—C5—H5A	120.6
O4—S1—O3	104.2 (3)	C4—C5—H5A	120.6
O2—S1—O3	102.7 (2)	C5—C4—C3	120.0 (4)
O1—S1—O3	108.7 (2)	C5—C4—H4A	120.0
C1—N1—C5	122.9 (4)	C3—C4—H4A	120.0
C1—N1—H1B	118.5	C1—C2—C3	119.5 (4)
C5—N1—H1B	118.5	C1—C2—H2A	120.2
C4—C3—C2	118.6 (4)	C3—C2—H2A	120.2
C4—C3—C6	123.0 (4)	S1—O3—H3	109.5
C2—C3—C6	118.5 (4)	C6—C7—H7A	109.5
N1—C1—C2	120.1 (4)	C6—C7—H7B	109.5
N1—C1—H1A	120.0	H7A—C7—H7B	109.5
C2—C1—H1A	120.0	C6—C7—H7C	109.5
O5—C6—C7	123.8 (5)	H7A—C7—H7C	109.5

O5—C6—C3	118.8 (5)	H7B—C7—H7C	109.5
C5—N1—C1—C2	-0.4 (7)	N1—C5—C4—C3	-0.1 (6)
C4—C3—C6—O5	157.9 (4)	C2—C3—C4—C5	-0.8 (6)
C2—C3—C6—O5	-21.8 (6)	C6—C3—C4—C5	179.5 (4)
C4—C3—C6—C7	-21.9 (6)	N1—C1—C2—C3	-0.5 (7)
C2—C3—C6—C7	158.4 (5)	C4—C3—C2—C1	1.1 (6)
C1—N1—C5—C4	0.7 (7)	C6—C3—C2—C1	-179.2 (4)

Hydrogen-bond geometry (Å, °)

<i>D—H...A</i>	<i>D—H</i>	<i>H...A</i>	<i>D...A</i>	<i>D—H...A</i>
N1—H1B...O1 ⁱ	0.86	1.92	2.772 (5)	172
O3—H3...O2 ⁱⁱ	0.82	1.76	2.565 (5)	166

Symmetry codes: (i) $-x+2, y+1/2, -z+1/2$; (ii) $x+1, y, z$.