Acta Crystallographica Section E
Structure Reports Online
ISSN 1600-5368

3-Aminopyridinium picrate

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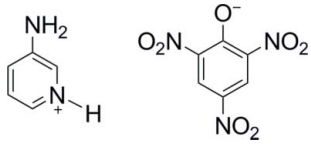
Received 5 September 2010; accepted 16 September 2010

Key indicators: single-crystal X-ray study; T = 297 K; mean σ(C–C) = 0.004 Å; R factor = 0.060; wR factor = 0.162; data-to-parameter ratio = 12.9.

During the formation of the title compound, $C_5H_7N_2^+\cdot C_6H_2N_3O_7^-$, a phenolic proton is transferred to the pyridine N atom. In the crystal structure, the ions are linked by intermolecular N—H···O and N—H···(O,O) hydrogen bonds into layers running parallel to (100). These layers are connected by weak π–π stacking interactions between symmetry-related pyridine and picric benzene rings with a centroid–centroid distance of 3.758 (2) Å, forming a three-dimensional network.

Related literature

For applications of picric acid derivatives, see: Pascard *et al.* (1982); Pearson *et al.* (2007); Shakir *et al.* (2009). For a related structure, see: Harrison *et al.* (2007).



Experimental

Crystal data

$C_5H_7N_2^+\cdot C_6H_2N_3O_7^-$
M_r = 323.23
Monoclinic, $P2_1/n$
a = 8.2174 (8) Å
b = 13.5842 (13) Å
c = 11.8218 (12) Å
β = 102.117 (2)°
V = 1290.2 (2) Å^3
Z = 4
Mo $K\alpha$ radiation
μ = 0.14 mm^{-1}
T = 297 K
0.45 × 0.05 × 0.02 mm

Data collection

Bruker SMART CCD diffractometer
Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996)
T_{min} = 0.939, T_{max} = 0.997
14192 measured reflections
2804 independent reflections
1391 reflections with $I > 2\sigma(I)$
R_{int} = 0.072

Refinement

$R[F^2 > 2\sigma(F^2)]$ = 0.060
$wR(F^2)$ = 0.162
S = 1.03
2804 reflections
217 parameters
H atoms treated by a mixture of independent and constrained refinement
$\Delta\rho_{max}$ = 0.34 e Å^{-3}
$\Delta\rho_{min}$ = −0.24 e Å^{-3}

Table 1
Hydrogen-bond geometry (Å, °).

D—H···A	D—H	H···A	D···A	D—H···A
N5—H5···O7	0.91 (3)	1.79 (3)	2.607 (3)	148 (3)
N5—H5···O6	0.91 (3)	2.46 (3)	3.179 (4)	136 (3)
N4—H4B···O6^{i}	0.86 (4)	2.48 (4)	3.119 (4)	132 (3)
N4—H4A···O3^{ii}	0.85 (4)	2.44 (4)	3.172 (4)	145 (3)

Symmetry codes: (i) $-x+\frac{3}{2}, y+\frac{1}{2}, -z+\frac{3}{2}$; (ii) $-x+\frac{3}{2}, y+\frac{1}{2}, -z+\frac{1}{2}$.

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINT* (Bruker, 2001); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *PLATON* (Spek, 2009); software used to prepare material for publication: *SHELXTL* (Sheldrick, 2008).

Wuhan University of Science and Technology is thanked for supporting this study.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: LH5128).

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supporting information

Acta Cryst. (2010). E**66**, o2680 [doi:10.1107/S1600536810037190]

3-Aminopyridinium picrate

Yan-jun Li

S1. Comment

Picric acid has been used in the characterization of organic bases because of the ease of crystallization and hence purification when picrate derivatives are produced (Pascard *et al.*, 1982; Pearson *et al.*, 2007; Harrison *et al.*, 2007; Shakir *et al.*, 2009). Here, we report the crystal structure of the title salt.

In the title compound, a hydrogen atom has been transferred from the picric acid molecule to the nitrogen atom of the pyridine ring and hence a 1:1 organic is formed salt (Fig.1). In the picric acid molecule, the geometric parameters of C6—O7 = 1.236 (3)Å and C1—C6—C5 = 112.0 (2)° confirm the transfer of the proton.

In the crystal structure, the molecular components are linked into a two dimensional zigzag-like layers (Fig.2) running parallel to (110) by intermolecular N—H···O hydrogen bonds (Table 1). These adjacent (100) layers are linked by weak π–π interaction between symmetry related pyridine and picric benzene rings (centroid-to-centroid distance = 3.758 (2) Å, symmetry code: 2 - *x*, 1 - *y*, 1 - *z*) into a three-dimensional network.

S2. Experimental

Picric acid (0.69 g, 3 mmol) and 3-aminopyridine (0.28 g, 3 mmol) were mixed in 10 ml ethanol. The mixture was kept at room temperature for two weeks. Yellow needeles suitable for single-crystal X-ray diffraction were obtained at the bottom of the vessel.

S3. Refinement

The carbon-bound hydrogen atoms were placed in ideal positions with C—H = 0.93Å and U_{iso}(H) = 1.2U$_{eq}$(C). The nitrogen-bound H atoms were located in a difference map and refined with U_{iso}(H) = 0.092Å^2.

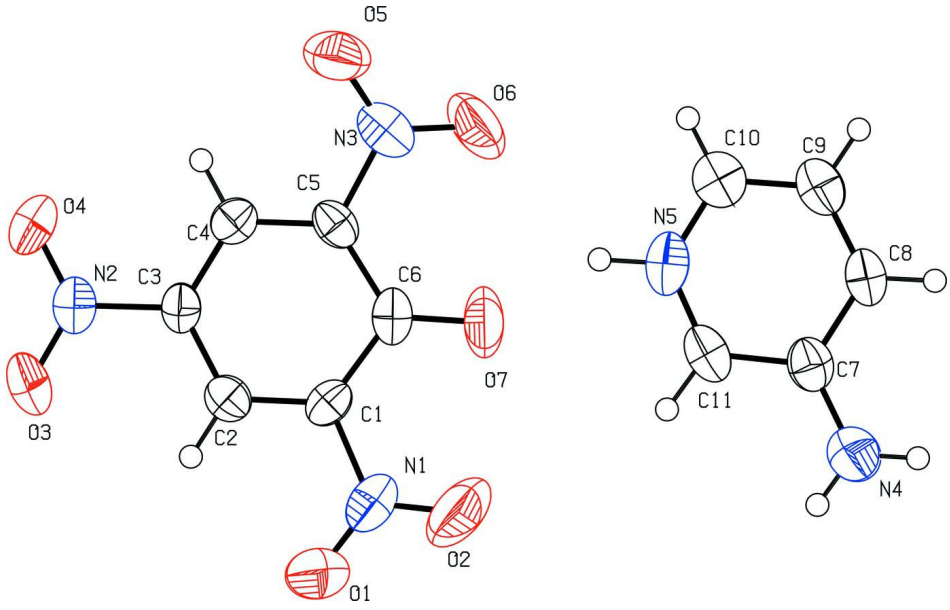

Figure 1

The asymmetric unit of the title compound with displacement ellipsoids drawn at the 30% probability level.

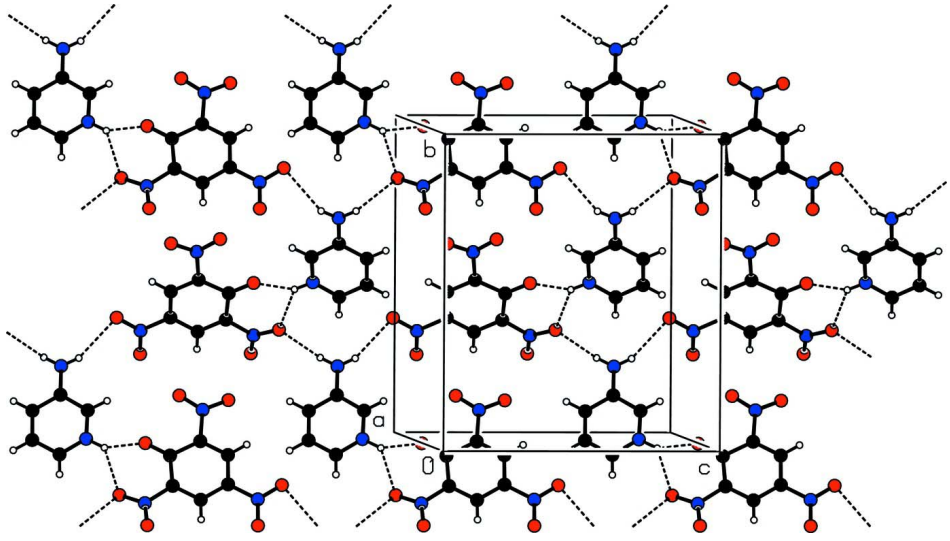

Figure 2

Part of the crystal structure with hydrogen bonds shown as dashed lines. For the sake of clarity, the H atoms not involved in the hydrogen-bonds pattern have been omitted.

3-Aminopyridinium picrate

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$C_5H_7N_2^+·C_6H_2N_3O_7^-$
M_r = 323.23
Monoclinic, $P2_1/n$
Hall symbol: -P 2yn
a = 8.2174 (8) Å
b = 13.5842 (13) Å
c = 11.8218 (12) Å
β = 102.117 (2)°
V = 1290.2 (2) Å^3
Z = 4
F(000) = 664
D_x = 1.664 Mg m^{-3}

Mo $K\alpha$ radiation, λ = 0.71073 Å
Cell parameters from 1100 reflections
θ = 2.3–20.4°
μ = 0.14 mm^{-1}
T = 297 K
Needle, yellow
0.45 × 0.05 × 0.02 mm

Data collection

Bruker SMART CCD diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
φ and ω scans
Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996)
T_{min} = 0.939, T_{max} = 0.997
14192 measured reflections
2804 independent reflections
1391 reflections with $I > 2\sigma(I)$
R_{int} = 0.072
θ_{max} = 27.0°, θ_{min} = 2.3°
h = −10→10
k = −17→17
l = −15→15

Refinement

Refinement on F^2
Least-squares matrix: full
$R[F^2 > 2\sigma(F^2)]$ = 0.060
$wR(F^2)$ = 0.162
S = 1.03
2804 reflections
217 parameters
0 restraints
Primary atom site location: structure-invariant direct methods
Secondary atom site location: difference Fourier map
Hydrogen site location: inferred from neighbouring sites
H atoms treated by a mixture of independent and constrained refinement
$w = 1/[\sigma^2(F_o^2) + (0.0663P)^2 + 0.0975P]$ where $P = (F_o^2 + 2F_c^2)/3$
$(\Delta/\sigma)_{max} < 0.001$
$\Delta\rho_{max}$ = 0.34 e Å^{-3}
$\Delta\rho_{min}$ = −0.24 e Å^{-3}

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 (Å²)

	x	y	z	U_{iso}*/U_{eq}
C1	0.8262 (3)	0.49045 (18)	0.2590 (2)	0.0390 (7)
C2	0.8816 (3)	0.45022 (19)	0.1686 (2)	0.0401 (7)
H2	0.8614	0.4819	0.0973	0.048*
C3	0.9673 (3)	0.36279 (18)	0.1827 (2)	0.0378 (7)
C4	1.0039 (3)	0.31637 (19)	0.2881 (2)	0.0388 (7)
H4	1.0636	0.2577	0.2967	0.047*
C5	0.9518 (3)	0.35703 (19)	0.3805 (2)	0.0388 (7)
C6	0.8524 (4)	0.4466 (2)	0.3728 (2)	0.0428 (7)
C7	0.5593 (4)	0.6203 (2)	0.7181 (2)	0.0449 (7)
C8	0.5870 (4)	0.5718 (2)	0.8241 (2)	0.0474 (8)
H8	0.5482	0.5994	0.8855	0.057*
C9	0.6702 (4)	0.4844 (2)	0.8392 (3)	0.0532 (8)

H9	0.6866	0.4527	0.9104	0.064*
C10	0.7300 (4)	0.4426 (2)	0.7507 (3)	0.0531 (8)
H10	0.7872	0.3831	0.7605	0.064*
C11	0.6221 (4)	0.5750 (2)	0.6304 (2)	0.0470 (8)
H11	0.6070	0.6043	0.5578	0.056*
N1	0.7366 (3)	0.58418 (19)	0.2352 (3)	0.0562 (7)
N2	1.0162 (3)	0.31865 (19)	0.0837 (2)	0.0498 (7)
N3	0.9978 (4)	0.3051 (2)	0.4890 (2)	0.0569 (7)
N4	0.4760 (4)	0.7061 (2)	0.7014 (3)	0.0684 (9)
N5	0.7032 (3)	0.4902 (2)	0.6502 (2)	0.0511 (7)
O1	0.6984 (4)	0.61237 (17)	0.1355 (3)	0.0962 (10)
O2	0.6942 (4)	0.6267 (2)	0.3121 (2)	0.1179 (12)
O3	0.9816 (3)	0.36138 (17)	−0.00931 (18)	0.0755 (8)
O4	1.0901 (3)	0.24022 (17)	0.09581 (19)	0.0736 (7)
O5	1.1054 (3)	0.24169 (18)	0.49917 (19)	0.0764 (8)
O6	0.9316 (3)	0.3268 (2)	0.5696 (2)	0.0916 (9)
O7	0.7949 (3)	0.48236 (16)	0.45224 (19)	0.0770 (8)
H4A	0.455 (5)	0.730 (3)	0.634 (3)	0.092*
H4B	0.436 (5)	0.731 (3)	0.757 (3)	0.092*
H5	0.749 (4)	0.465 (2)	0.592 (3)	0.092*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0361 (16)	0.0317 (15)	0.0490 (17)	0.0008 (12)	0.0087 (14)	−0.0052 (13)
C2	0.0394 (17)	0.0427 (16)	0.0376 (16)	−0.0036 (14)	0.0069 (13)	0.0015 (13)
C3	0.0414 (17)	0.0381 (16)	0.0369 (16)	−0.0046 (13)	0.0147 (13)	−0.0068 (12)
C4	0.0371 (16)	0.0341 (15)	0.0453 (16)	−0.0027 (13)	0.0085 (13)	−0.0026 (13)
C5	0.0419 (17)	0.0425 (16)	0.0319 (15)	−0.0109 (13)	0.0075 (13)	0.0019 (12)
C6	0.0417 (18)	0.0483 (17)	0.0422 (17)	−0.0090 (14)	0.0171 (14)	−0.0127 (14)
C7	0.0475 (18)	0.0477 (18)	0.0402 (17)	−0.0098 (15)	0.0106 (14)	−0.0062 (14)
C8	0.0508 (19)	0.0570 (19)	0.0380 (17)	−0.0061 (16)	0.0178 (14)	−0.0072 (14)
C9	0.056 (2)	0.065 (2)	0.0385 (17)	−0.0058 (17)	0.0098 (16)	−0.0019 (15)
C10	0.049 (2)	0.057 (2)	0.0526 (19)	−0.0070 (15)	0.0094 (16)	−0.0019 (16)
C11	0.0470 (19)	0.061 (2)	0.0333 (16)	−0.0152 (16)	0.0095 (14)	−0.0029 (14)
N1	0.0469 (16)	0.0525 (17)	0.0693 (19)	0.0071 (13)	0.0129 (15)	−0.0143 (15)
N2	0.0528 (17)	0.0548 (17)	0.0453 (16)	−0.0010 (13)	0.0183 (13)	−0.0087 (13)
N3	0.0603 (19)	0.0655 (18)	0.0432 (16)	−0.0104 (16)	0.0070 (14)	0.0083 (14)
N4	0.095 (2)	0.0606 (19)	0.0508 (18)	0.0098 (17)	0.0180 (17)	0.0020 (15)
N5	0.0461 (17)	0.0593 (17)	0.0514 (18)	−0.0099 (13)	0.0184 (13)	−0.0191 (14)
O1	0.127 (2)	0.0793 (18)	0.094 (2)	0.0456 (17)	0.0484 (19)	0.0347 (15)
O2	0.144 (3)	0.109 (2)	0.087 (2)	0.074 (2)	−0.0071 (19)	−0.0394 (17)
O3	0.100 (2)	0.0951 (18)	0.0367 (12)	0.0179 (15)	0.0255 (12)	0.0006 (12)
O4	0.0958 (19)	0.0592 (14)	0.0733 (16)	0.0264 (14)	0.0347 (14)	−0.0095 (12)
O5	0.098 (2)	0.0659 (16)	0.0572 (15)	0.0144 (15)	−0.0020 (14)	0.0129 (12)
O6	0.094 (2)	0.136 (2)	0.0549 (15)	0.0175 (17)	0.0390 (15)	0.0328 (15)
O7	0.106 (2)	0.0784 (16)	0.0612 (15)	0.0045 (14)	0.0509 (15)	−0.0147 (12)

Geometric parameters (Å, º)

C1—C2	1.361 (3)	C8—H8	0.9300
C1—C6	1.446 (4)	C9—C10	1.369 (4)
C1—N1	1.468 (3)	C9—H9	0.9300
C2—C3	1.373 (3)	C10—N5	1.329 (4)
C2—H2	0.9300	C10—H10	0.9300
C3—C4	1.372 (4)	C11—N5	1.328 (4)
C3—N2	1.445 (3)	C11—H11	0.9300
C4—C5	1.371 (3)	N1—O2	1.190 (3)
C4—H4	0.9300	N1—O1	1.216 (3)
C5—N3	1.443 (3)	N2—O4	1.220 (3)
C5—C6	1.457 (4)	N2—O3	1.223 (3)
C6—O7	1.236 (3)	N3—O5	1.222 (3)
C7—N4	1.345 (4)	N3—O6	1.229 (3)
C7—C8	1.392 (4)	N4—H4A	0.85 (4)
C7—C11	1.394 (4)	N4—H4B	0.86 (4)
C8—C9	1.364 (4)	N5—H5	0.91 (3)
C2—C1—C6	123.8 (2)	C8—C9—C10	120.8 (3)
C2—C1—N1	115.8 (3)	C8—C9—H9	119.6
C6—C1—N1	120.4 (3)	C10—C9—H9	119.6
C1—C2—C3	120.0 (3)	N5—C10—C9	117.5 (3)
C1—C2—H2	120.0	N5—C10—H10	121.2
C3—C2—H2	120.0	C9—C10—H10	121.2
C4—C3—C2	121.1 (2)	N5—C11—C7	120.2 (3)
C4—C3—N2	120.0 (2)	N5—C11—H11	119.9
C2—C3—N2	118.9 (2)	C7—C11—H11	119.9
C5—C4—C3	119.5 (3)	O2—N1—O1	122.0 (3)
C5—C4—H4	120.2	O2—N1—C1	119.3 (3)
C3—C4—H4	120.2	O1—N1—C1	118.4 (3)
C4—C5—N3	116.4 (3)	O4—N2—O3	122.4 (2)
C4—C5—C6	123.4 (2)	O4—N2—C3	118.9 (3)
N3—C5—C6	120.2 (3)	O3—N2—C3	118.6 (3)
O7—C6—C1	122.6 (3)	O5—N3—O6	121.5 (3)
O7—C6—C5	125.4 (3)	O5—N3—C5	118.8 (3)
C1—C6—C5	112.0 (2)	O6—N3—C5	119.7 (3)
N4—C7—C8	121.6 (3)	C7—N4—H4A	118 (3)
N4—C7—C11	122.0 (3)	C7—N4—H4B	119 (3)
C8—C7—C11	116.4 (3)	H4A—N4—H4B	122 (4)
C9—C8—C7	120.8 (3)	C11—N5—C10	124.2 (3)
C9—C8—H8	119.6	C11—N5—H5	117 (2)
C7—C8—H8	119.6	C10—N5—H5	118 (2)
C6—C1—C2—C3	0.3 (4)	C7—C8—C9—C10	0.6 (4)
N1—C1—C2—C3	179.8 (2)	C8—C9—C10—N5	−0.2 (4)
C1—C2—C3—C4	−2.4 (4)	N4—C7—C11—N5	−179.8 (3)
C1—C2—C3—N2	176.4 (2)	C8—C7—C11—N5	0.0 (4)

C2—C3—C4—C5	1.1 (4)	C2—C1—N1—O2	−175.1 (3)
N2—C3—C4—C5	−177.7 (2)	C6—C1—N1—O2	4.4 (4)
C3—C4—C5—N3	−178.8 (2)	C2—C1—N1—O1	9.8 (4)
C3—C4—C5—C6	2.4 (4)	C6—C1—N1—O1	−170.7 (3)
C2—C1—C6—O7	−176.9 (3)	C4—C3—N2—O4	−0.1 (4)
N1—C1—C6—O7	3.6 (4)	C2—C3—N2—O4	−178.9 (3)
C2—C1—C6—C5	2.8 (4)	C4—C3—N2—O3	179.8 (3)
N1—C1—C6—C5	−176.7 (2)	C2—C3—N2—O3	1.0 (4)
C4—C5—C6—O7	175.5 (3)	C4—C5—N3—O5	14.6 (4)
N3—C5—C6—O7	−3.2 (4)	C6—C5—N3—O5	−166.5 (3)
C4—C5—C6—C1	−4.1 (4)	C4—C5—N3—O6	−167.1 (3)
N3—C5—C6—C1	177.1 (2)	C6—C5—N3—O6	11.8 (4)
N4—C7—C8—C9	179.3 (3)	C7—C11—N5—C10	0.4 (4)
C11—C7—C8—C9	−0.5 (4)	C9—C10—N5—C11	−0.3 (4)

Hydrogen-bond geometry (Å, º)

D—H···*A*	*D*—H	H···*A*	*D*···*A*	*D*—H···*A*
N5—H5···O7	0.91 (3)	1.79 (3)	2.607 (3)	148 (3)
N5—H5···O6	0.91 (3)	2.46 (3)	3.179 (4)	136 (3)
N4—H4*B*···O6^{i}	0.86 (4)	2.48 (4)	3.119 (4)	132 (3)
N4—H4*A*···O3^{ii}	0.85 (4)	2.44 (4)	3.172 (4)	145 (3)

Symmetry codes: (i) −*x*+3/2, *y*+1/2, −*z*+3/2; (ii) −*x*+3/2, *y*+1/2, −*z*+1/2.