

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.049$
 $wR(F^2) = 0.125$
 $S = 0.91$
 2791 reflections

111 parameters
 H-atom parameters constrained
 $\Delta\rho_{\max} = 0.29 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.19 \text{ e } \text{\AA}^{-3}$

4-Methylanilinium nitrate

Nouredine Benali-Cherif,* Houda Boussekine, Zina Boutobba and Noureddine Dadda

Laboratoire des Structures, Propriétés et Interactions Inter Atomiques (LASPI²A), Centre Universitaire Abbes Laghrour-Khenchela, 40000. Khenchela, Algeria
 Correspondence e-mail: benalicherif@hotmail.com

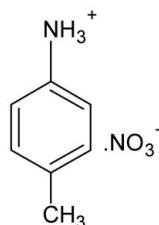
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Key indicators: single-crystal X-ray study; $T = 100 \text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.002 \text{ \AA}$; R factor = 0.049; wR factor = 0.125; data-to-parameter ratio = 25.1.

The asymmetric unit of the title compound, $\text{C}_7\text{H}_{10}\text{N}^+\cdot\text{NO}_3^-$, consists of a 4-methylanilinium cation protonated at the amino group and a nitrate anion. In the crystal, anions and cations are linked through $\text{N}-\text{H}\cdots\text{O}$ and $\text{N}-\text{H}\cdots(\text{O},\text{O})$ hydrogen bonds, building a corrugated layer structure parallel to (001).

Related literature

For related structures, see: Benali-Cherif, Kateb *et al.* (2007); Benali-Cherif, Allouche *et al.* (2007); Benali-Cherif, Boussekine *et al.* (2007); Asath Bahadur *et al.* (2007). For the biological effects of toluidine exposure in man, see: Kennedy *et al.* (1984).



Experimental

Crystal data

$\text{C}_7\text{H}_{10}\text{N}^+\cdot\text{NO}_3^-$
 $M_r = 170.17$
 Monoclinic, $P2_1/n$
 $a = 5.6725 (9) \text{ \AA}$
 $b = 8.5507 (8) \text{ \AA}$
 $c = 17.621 (2) \text{ \AA}$
 $\beta = 98.771 (2)^\circ$

 $V = 844.69 (18) \text{ \AA}^3$
 $Z = 4$

 Mo $K\alpha$ radiation

 $\mu = 0.11 \text{ mm}^{-1}$
 $T = 100 \text{ K}$
 $0.2 \times 0.15 \times 0.1 \text{ mm}$

Data collection

Nonius KappaCCD diffractometer
 Absorption correction: none
 24550 measured reflections

2791 independent reflections
 1228 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.089$

Table 1
 Hydrogen-bond geometry (\AA , $^\circ$).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
N1—H1B \cdots O3	0.89	1.93	2.8032 (17)	167
N1—H1A \cdots O2 ⁱ	0.89	1.93	2.8208 (18)	177
N1—H1C \cdots O3 ⁱⁱ	0.89	2.11	2.9461 (17)	157
N1—H1C \cdots O2 ⁱⁱ	0.89	2.46	3.1726 (18)	138

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

Data collection: *KappaCCD* (Nonius, 1998); cell refinement: *DENZO* and *SCALEPACK* (Otwinowski & Minor, 1997); data reduction: *DENZO* and *SCALEPACK*; program(s) used to solve structure: *SIR2004* (Burla *et al.*, 2005); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3* (Farrugia, 1997) and *CAMERON* (Pearce *et al.*, 2000); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: DN2484).

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4-Methylanilinium nitrate

N. Benali-Cherif, H. Boussekine, Z. Boutobba and N. Dadda

Comment

p-toluidine is an organic benzene derivative with a methyl substituent and an amino group, the name is derived from toluene and aniline. Its physical appearance is that of white lustrous plates or leaflets with an amine odour. *p*-toluidine can cause anoxia (due to formation of methemoglobin) and hematuria in man. The substance irritates the eyes and the skin and may cause effects on the blood, bladder and kidneys, resulting in tissue lesions and formation of methamoglobin (Kennedy *et al.*, 1984). The crystal structure of *p*-methylanilinium nitrate, (I), was determined as part of our investigations on the structural characteristics of organic-inorganic layered compounds and an ongoing study on *D*—*H*···*A* hydrogen-bonding in systems of hybrid materials including anilinium derivatives such as, 3-hydroxyanilinium hydrogensulfate (Benali-Cherif, Kateb *et al.*, 2007), *o*-methylanilinium nitrate (Benali-Cherif, Boussekine *et al.*, 2007), 2-carboxyanilinium dihydrogenphosphate (Benali-Cherif, Allouche *et al.*, 2007) and 2-carboxyanilinium nitrate (Bahadur *et al.*, 2007).

The asymmetric unit of (I) contains a monoprotonated *p*-methylanilinium cation and nitrate anion linked through N—H···O hydrogen bond (Figure 1). Intra atomic bond distances and angles confirm the monoprotonation of the organic entity. There are differences in the N—O distances of nitrate anion N2—O2, N2—O3 (1.260 (2) Å, 1.276 (2) Å) are longer than N2—O1 (1.232 (2) Å), this is due to the fact that only the O2 and O3 atoms are involved in hydrogen bonds of types N—H···O. (Table 1). The structure of ($C_7H_{10}N^+$. NO_3^-) is composed of cationic ($C_7H_{10}N^+$) and anionic (NO_3^-) linked through N—H···O hydrogen bonds and building up a corrugated layers parallel to the (0 0 1) plane (Table 1, Figure 2).

Experimental

Single crystals of the title compound are prepared by slow evaporation at room temperature of an aqueous solution of *p*-methylaniline (C_7H_9N) and nitric acid in the stoichiometric ratio 1:1.

Refinement

All H atoms attached to C atoms and N atom were fixed geometrically and treated as riding with C—H = 0.96 Å (methyl) or 0.93 Å (aromatic) and N—H = 0.89 Å with $U_{iso}(H) = 1.2U_{eq}(\text{aromatic})$ or $U_{iso}(H) = 1.5U_{eq}(\text{methyl}, \text{N})$.

Figures

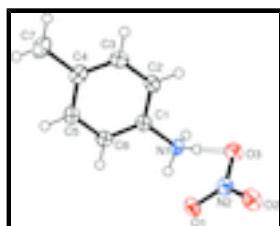


Fig. 1. Molecular view of compound I with the atom labeling scheme. Displacement ellipsoids are drawn at the 50% probability level. H atoms are represented as small spheres of arbitrary radii. Hydrogen bond is shown as dashed line.

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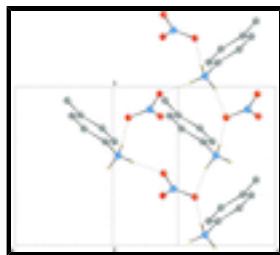


Fig. 2. Partial packing view of the hydrogen-bonding network.

4-Methylanilinium nitrate

Crystal data

$C_7H_{10}N^+\cdot NO_3^-$	$F_{000} = 360$
$M_r = 170.17$	$D_x = 1.338 \text{ Mg m}^{-3}$
Monoclinic, $P2_1/n$	$Mo K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$
Hall symbol: -P 2yn	Cell parameters from 24550 reflections
$a = 5.6725 (9) \text{ \AA}$	$\theta = 2.7\text{--}31.5^\circ$
$b = 8.5507 (8) \text{ \AA}$	$\mu = 0.11 \text{ mm}^{-1}$
$c = 17.621 (2) \text{ \AA}$	$T = 100 \text{ K}$
$\beta = 98.771 (2)^\circ$	Prism, brown
$V = 844.69 (18) \text{ \AA}^3$	$0.2 \times 0.15 \times 0.1 \text{ mm}$
$Z = 4$	

Data collection

Nonius KappaCCD diffractometer	1228 reflections with $I > 2\sigma(I)$
Radiation source: fine-focus sealed tube	$R_{\text{int}} = 0.089$
Monochromator: graphite	$\theta_{\text{max}} = 31.5^\circ$
$T = 100 \text{ K}$	$\theta_{\text{min}} = 2.7^\circ$
$\omega\text{--}\theta$ scans	$h = -8 \rightarrow 5$
Absorption correction: none	$k = -12 \rightarrow 12$
24550 measured reflections	$l = -25 \rightarrow 25$
2791 independent reflections	

Refinement

Refinement on F^2	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.049$	H-atom parameters constrained
$wR(F^2) = 0.125$	$w = 1/[\sigma^2(F_o^2) + (0.0589P)^2]$
$S = 0.91$	where $P = (F_o^2 + 2F_c^2)/3$
2791 reflections	$(\Delta/\sigma)_{\text{max}} = 0.001$
111 parameters	$\Delta\rho_{\text{max}} = 0.29 \text{ e \AA}^{-3}$
	$\Delta\rho_{\text{min}} = -0.19 \text{ e \AA}^{-3}$

Primary atom site location: structure-invariant direct methods Extinction correction: none

Special details

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

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.4327 (3)	0.15225 (17)	0.90563 (9)	0.0212 (4)
C2	0.2258 (3)	0.23416 (18)	0.91199 (10)	0.0251 (4)
H2	0.0979	0.2342	0.8721	0.030*
C3	0.2140 (3)	0.31609 (19)	0.97933 (10)	0.0280 (4)
H3	0.0752	0.3705	0.9843	0.034*
C4	0.4025 (3)	0.31936 (17)	1.03941 (10)	0.0261 (4)
C5	0.6101 (3)	0.23714 (18)	1.03075 (9)	0.0265 (4)
H5	0.7396	0.2386	1.0701	0.032*
C6	0.6249 (3)	0.15283 (17)	0.96364 (9)	0.0239 (4)
H6	0.7629	0.0979	0.9583	0.029*
C7	0.3867 (4)	0.4121 (2)	1.11156 (10)	0.0347 (5)
H7A	0.4333	0.5184	1.1044	0.052*
H7B	0.4911	0.3670	1.1539	0.052*
H7C	0.2257	0.4099	1.1222	0.052*
N1	0.4466 (2)	0.06544 (14)	0.83447 (7)	0.0236 (3)
H1A	0.5927	0.0263	0.8361	0.035*
H1B	0.4147	0.1296	0.7945	0.035*
H1C	0.3409	-0.0122	0.8298	0.035*
N2	0.5736 (3)	0.35591 (15)	0.71667 (8)	0.0249 (3)
O1	0.7537 (2)	0.31274 (13)	0.75970 (7)	0.0311 (3)
O2	0.5852 (2)	0.45260 (13)	0.66347 (7)	0.0339 (3)
O3	0.3676 (2)	0.30432 (13)	0.72457 (7)	0.0295 (3)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0282 (9)	0.0117 (7)	0.0233 (9)	-0.0041 (7)	0.0031 (7)	0.0006 (6)
C2	0.0245 (9)	0.0180 (8)	0.0312 (10)	-0.0018 (7)	-0.0012 (7)	0.0002 (7)
C3	0.0290 (10)	0.0182 (8)	0.0373 (11)	0.0018 (7)	0.0066 (8)	-0.0007 (7)
C4	0.0381 (10)	0.0130 (7)	0.0285 (10)	-0.0053 (7)	0.0095 (8)	0.0016 (7)
C5	0.0333 (10)	0.0207 (8)	0.0241 (10)	-0.0042 (7)	0.0001 (8)	0.0029 (7)

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C6	0.0261 (9)	0.0167 (8)	0.0291 (10)	0.0000 (7)	0.0051 (7)	0.0036 (7)
C7	0.0524 (12)	0.0222 (8)	0.0305 (11)	-0.0023 (8)	0.0097 (9)	-0.0023 (7)
N1	0.0277 (8)	0.0166 (6)	0.0260 (8)	-0.0014 (6)	0.0022 (6)	0.0002 (6)
N2	0.0290 (8)	0.0169 (7)	0.0279 (8)	0.0011 (6)	0.0018 (7)	-0.0022 (6)
O1	0.0276 (7)	0.0305 (7)	0.0326 (7)	0.0051 (6)	-0.0038 (6)	0.0023 (6)
O2	0.0354 (7)	0.0252 (6)	0.0397 (8)	-0.0020 (6)	0.0008 (6)	0.0139 (6)
O3	0.0277 (7)	0.0267 (6)	0.0338 (7)	-0.0015 (5)	0.0041 (5)	0.0052 (5)

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

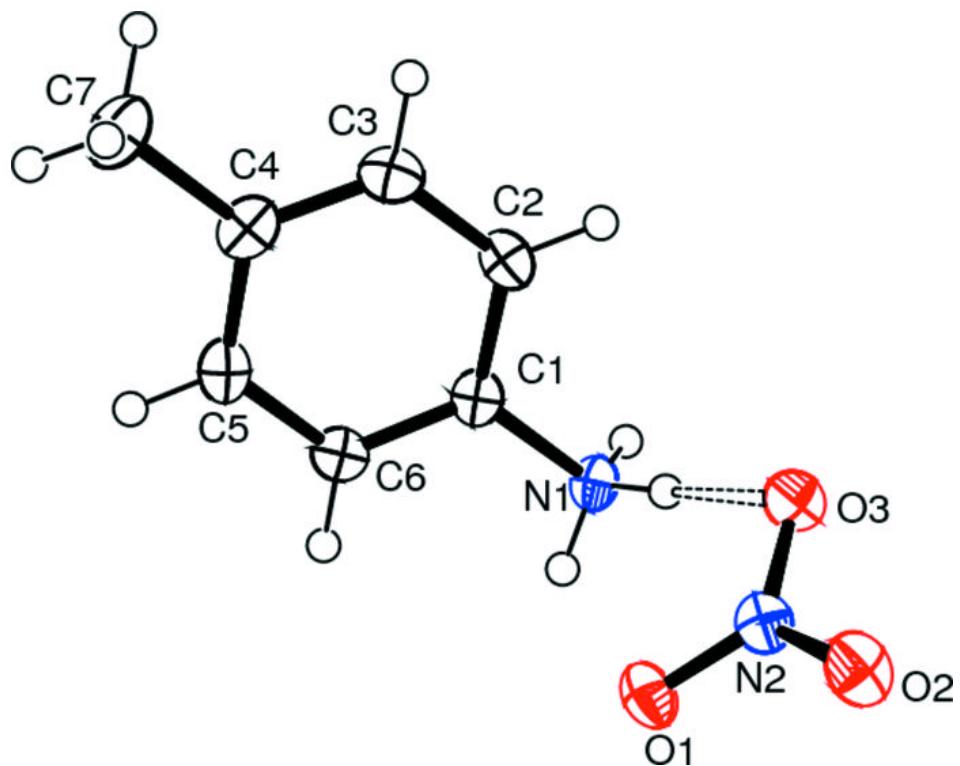
C1—C6	1.376 (2)	C6—H6	0.9300
C1—C2	1.386 (2)	C7—H7A	0.9600
C1—N1	1.470 (2)	C7—H7B	0.9600
C2—C3	1.388 (2)	C7—H7C	0.9600
C2—H2	0.9300	N1—H1A	0.8900
C3—C4	1.385 (2)	N1—H1B	0.8900
C3—H3	0.9300	N1—H1C	0.8900
C4—C5	1.399 (2)	N2—O1	1.2325 (17)
C4—C7	1.513 (2)	N2—O2	1.2592 (16)
C5—C6	1.398 (2)	N2—O3	1.2759 (17)
C5—H5	0.9300		
C6—C1—C2	121.53 (15)	C5—C6—H6	120.4
C6—C1—N1	119.74 (14)	C4—C7—H7A	109.5
C2—C1—N1	118.73 (14)	C4—C7—H7B	109.5
C1—C2—C3	118.41 (16)	H7A—C7—H7B	109.5
C1—C2—H2	120.8	C4—C7—H7C	109.5
C3—C2—H2	120.8	H7A—C7—H7C	109.5
C4—C3—C2	122.09 (16)	H7B—C7—H7C	109.5
C4—C3—H3	119.0	C1—N1—H1A	109.5
C2—C3—H3	119.0	C1—N1—H1B	109.5
C3—C4—C5	118.08 (16)	H1A—N1—H1B	109.5
C3—C4—C7	121.01 (16)	C1—N1—H1C	109.5
C5—C4—C7	120.90 (17)	H1A—N1—H1C	109.5
C6—C5—C4	120.75 (16)	H1B—N1—H1C	109.5
C6—C5—H5	119.6	O1—N2—O2	121.47 (14)
C4—C5—H5	119.6	O1—N2—O3	121.07 (14)
C1—C6—C5	119.14 (16)	O2—N2—O3	117.45 (14)
C1—C6—H6	120.4		

Hydrogen-bond geometry (\AA , $^\circ$)

$D\cdots H$	$H\cdots A$	$D\cdots A$	$D—H\cdots A$
N1—H1B···O3	0.89	1.93	2.8032 (17)
N1—H1A···O2 ⁱ	0.89	1.93	2.8208 (18)
N1—H1C···O3 ⁱⁱ	0.89	2.11	2.9461 (17)
N1—H1C···O2 ⁱⁱ	0.89	2.46	3.1726 (18)

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

Fig. 1



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Fig. 2

