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2-Methylbenzimidazolium nitrate

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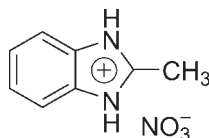
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Key indicators: single-crystal X-ray study; $T = 298$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; R factor = 0.045; wR factor = 0.135; data-to-parameter ratio = 13.2.

In the title compound, $\text{C}_8\text{H}_9\text{N}_2^+\cdot\text{NO}_3^-$, intermolecular $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds join the molecules into a chain extending along the b axis.

Related literature

For the applications of related benzimidazole compounds, see: Wright (1951); El-masry *et al.* (2000); Gümüş *et al.* (2003).



Experimental

Crystal data

 $\text{C}_8\text{H}_9\text{N}_2^+\cdot\text{NO}_3^-$ $M_r = 195.18$ Monoclinic, $P2_1/c$ $a = 7.711$ (4) Å $b = 15.127$ (7) Å $c = 8.270$ (4) Å $\beta = 99.398$ (7)° $V = 951.7$ (8) Å³ $Z = 4$ Mo $K\alpha$ radiation $\mu = 0.11$ mm⁻¹ $T = 298$ K

0.18 × 0.16 × 0.12 mm

Data collection

Bruker SMART APEX diffractometer
Absorption correction: multi-scan (SADABS; Bruker, 2005)
 $T_{\min} = 0.981$, $T_{\max} = 0.987$

4774 measured reflections
1685 independent reflections
1319 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.027$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.045$
 $wR(F^2) = 0.135$
 $S = 1.07$
1685 reflections

128 parameters
H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.22$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.21$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{N2}-\text{H2}\cdots\text{O1}^{\text{i}}$	0.86	2.03	2.855 (3)	162
$\text{N3}-\text{H3}\cdots\text{O2}^{\text{ii}}$	0.86	1.93	2.775 (2)	166

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

Data collection: SMART (Bruker, 2005); cell refinement: SAINT (Bruker, 2005); data reduction: SAINT; 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: SHELXL97.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: GK2255).

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supporting information

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2-Methylbenzimidazolium nitrate

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S1. Comment

Benzimidazole and its derivatives have found practical applications in a number of fields (Wright, 1951). This ring system is present in numerous antiparasitic, antihelmintic and anti-inflammatory drugs (El-masry *et al.*, 2000). The complexes of transition metals with benzimidazole and related ligands have been extensively studied as models of some important biological molecules (Gümüş *et al.*, 2003). During our search to find new benzimidazole-metal complexes 2-methylbenzimidazole nitrate was unintentionally obtained.

Herein, we report the structure of the title compound, C₈H₉N₃O₃ (Fig 1). The crystal structure showed that intermolecular N—H···O hydrogen bonds link the molecules into a 1D polymeric structure (Fig. 2).

S2. Experimental

A mixture of *o*-phenylenediamine (1.08 g, 10 mmol) and anhydrous sodium acetate (2.46 g, 30 mmol) were dissolved in 100 mL 5% hydrochloric acid. After stirring for 2 h under reflux, the solution was cooled to room temperature. Then the solution was treated with ammonia solution to pH 9-10 and an orange precipitate was formed. The precipitate was filtered and washed with water. 2-methylbenzimidazolium chloride was gained in 27.32% yield. The compound 2-methylbenzimidazole nitrate was obtained in 35% yield when the 2-methylbenzimidazolium chloride (0.46 g, 2.73 mmol) was reacted with Cr(NO₃)₃·9H₂O (1.01 g, 2.54 mmol) in ethanol under reflux. The crystals suitable for X-ray diffraction analysis were obtained by recrystallization from ethanol.

S3. Refinement

All H atoms were located in difference maps. H atoms bonded to C atoms were then treated as riding atoms in geometrically idealized positions, with C—H distances of 0.93 (aromatic), 0.96 (CH₃—H) and 0.86 (N—H) Å, and with U_{iso}(H) = kU_{eq}(C), where k is 1.5 for the methyl group and 1.2 for all the other H atoms.

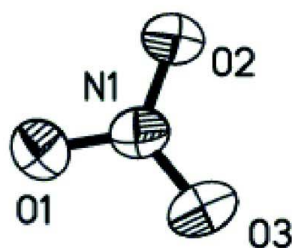
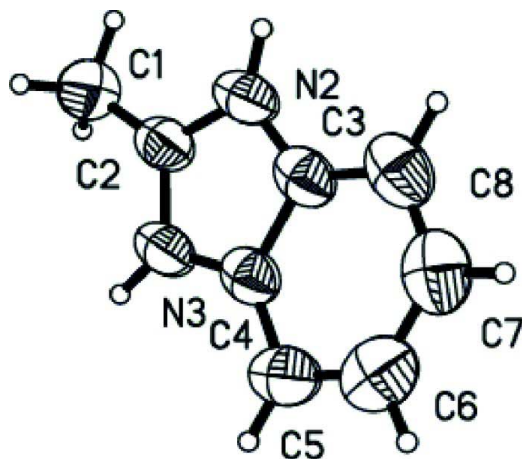


Figure 1

The structure of the title compound showing 50% probability displacement.

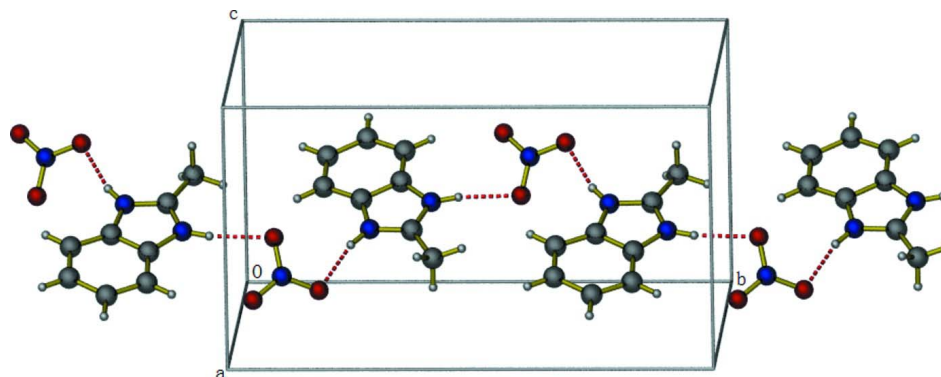


Figure 2

The supramolecular chain of the title compound formed via N—H...O hydrogen bonds.

2-methylbenzimidazolium nitrate

Crystal data

$C_8H_9N_2^+ \cdot NO_3^-$

$M_r = 195.18$

Monoclinic, $P2_1/c$

Hall symbol: $-P\ 2ybc$

$a = 7.711(4)\ \text{\AA}$

$b = 15.127(7)\ \text{\AA}$

$c = 8.270(4)\ \text{\AA}$

$\beta = 99.398(7)^\circ$

$V = 951.7(8)\ \text{\AA}^3$

$Z = 4$

$F(000) = 408$

$D_x = 1.362\ \text{Mg m}^{-3}$

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

Cell parameters from 2174 reflections

$\theta = 2.5\text{--}25.5^\circ$

$\mu = 0.11\ \text{mm}^{-1}$

$T = 298\ \text{K}$

Block, colorless

$0.18 \times 0.16 \times 0.12\ \text{mm}$

Data collection

Bruker SMART APEX
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

φ and ω scans

Absorption correction: multi-scan
(*SADABS*; Bruker, 2005)

$T_{\min} = 0.981$, $T_{\max} = 0.987$

4774 measured reflections

1685 independent reflections

1319 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.027$

$\theta_{\max} = 25.1^\circ$, $\theta_{\min} = 2.7^\circ$

$h = -9 \rightarrow 9$

$k = -11 \rightarrow 18$

$l = -9 \rightarrow 9$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.045$

$wR(F^2) = 0.135$

$S = 1.07$

1685 reflections

128 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-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0681P)^2 + 0.208P]$

where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} < 0.001$

$\Delta\rho_{\max} = 0.22\ \text{e \AA}^{-3}$

$\Delta\rho_{\min} = -0.21\ \text{e \AA}^{-3}$

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>	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.3641 (2)	0.06287 (10)	0.78462 (18)	0.0735 (5)
O2	0.33057 (19)	0.15524 (9)	0.97555 (17)	0.0643 (4)
O3	0.2248 (3)	0.02349 (11)	0.9773 (2)	0.0915 (6)
N1	0.3063 (2)	0.07987 (10)	0.9136 (2)	0.0538 (4)
N2	0.6679 (2)	0.12445 (11)	0.25067 (19)	0.0600 (5)
H2	0.6828	0.0687	0.2374	0.072*
N3	0.5577 (2)	0.24731 (11)	0.31726 (19)	0.0576 (5)
H3	0.4889	0.2842	0.3542	0.069*
C1	0.3941 (3)	0.11166 (15)	0.3743 (3)	0.0727 (6)
H1A	0.3921	0.0516	0.3368	0.109*
H1B	0.2839	0.1395	0.3331	0.109*
H1C	0.4132	0.1126	0.4920	0.109*
C2	0.5376 (3)	0.15990 (13)	0.3145 (2)	0.0569 (5)
C3	0.7769 (3)	0.19049 (13)	0.2083 (2)	0.0561 (5)
C4	0.7057 (3)	0.26993 (13)	0.2519 (2)	0.0542 (5)
C5	0.7834 (3)	0.35057 (14)	0.2303 (3)	0.0677 (6)
H5	0.7356	0.4035	0.2595	0.081*
C6	0.9355 (3)	0.34816 (19)	0.1630 (3)	0.0805 (7)
H6	0.9920	0.4009	0.1463	0.097*
C7	1.0070 (3)	0.2684 (2)	0.1191 (3)	0.0802 (7)
H7	1.1095	0.2695	0.0734	0.096*
C8	0.9302 (3)	0.18829 (18)	0.1416 (3)	0.0715 (7)
H8	0.9787	0.1353	0.1134	0.086*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.1006 (12)	0.0560 (9)	0.0718 (10)	0.0061 (8)	0.0370 (9)	-0.0048 (7)
O2	0.0865 (10)	0.0434 (8)	0.0651 (9)	-0.0085 (7)	0.0188 (7)	-0.0067 (6)
O3	0.1307 (16)	0.0554 (10)	0.0990 (13)	-0.0287 (10)	0.0499 (12)	0.0017 (8)
N1	0.0628 (10)	0.0406 (9)	0.0590 (10)	0.0020 (7)	0.0127 (8)	0.0042 (7)
N2	0.0735 (11)	0.0464 (9)	0.0564 (10)	0.0153 (8)	-0.0006 (8)	-0.0086 (7)
N3	0.0710 (11)	0.0476 (10)	0.0521 (9)	0.0150 (8)	0.0038 (8)	-0.0094 (7)
C1	0.0902 (16)	0.0612 (14)	0.0673 (13)	0.0017 (12)	0.0145 (12)	-0.0006 (11)
C2	0.0728 (13)	0.0487 (12)	0.0462 (10)	0.0126 (10)	0.0004 (9)	-0.0052 (8)
C3	0.0621 (12)	0.0575 (12)	0.0440 (10)	0.0131 (10)	-0.0051 (8)	-0.0086 (9)

C4	0.0619 (11)	0.0536 (11)	0.0433 (10)	0.0096 (9)	-0.0024 (8)	-0.0058 (8)
C5	0.0810 (15)	0.0539 (13)	0.0631 (13)	0.0037 (11)	-0.0033 (11)	-0.0019 (10)
C6	0.0806 (16)	0.0842 (18)	0.0716 (15)	-0.0133 (14)	-0.0030 (12)	0.0061 (13)
C7	0.0679 (14)	0.105 (2)	0.0662 (14)	0.0061 (14)	0.0070 (11)	-0.0032 (14)
C8	0.0685 (14)	0.0831 (17)	0.0593 (13)	0.0169 (13)	-0.0007 (11)	-0.0124 (11)

Geometric parameters (Å, °)

O1—N1	1.248 (2)	C1—H1C	0.9600
O2—N1	1.252 (2)	C3—C8	1.384 (3)
O3—N1	1.228 (2)	C3—C4	1.393 (3)
N2—C2	1.322 (3)	C4—C5	1.383 (3)
N2—C3	1.387 (3)	C5—C6	1.378 (4)
N2—H2	0.8600	C5—H5	0.9300
N3—C2	1.331 (3)	C6—C7	1.399 (4)
N3—C4	1.383 (3)	C6—H6	0.9300
N3—H3	0.8600	C7—C8	1.374 (4)
C1—C2	1.476 (3)	C7—H7	0.9300
C1—H1A	0.9600	C8—H8	0.9300
C1—H1B	0.9600		
O3—N1—O1	120.19 (17)	C8—C3—N2	132.5 (2)
O3—N1—O2	120.66 (17)	C8—C3—C4	121.5 (2)
O1—N1—O2	119.14 (16)	N2—C3—C4	105.96 (18)
C2—N2—C3	109.90 (17)	C5—C4—N3	132.13 (19)
C2—N2—H2	125.1	C5—C4—C3	122.0 (2)
C3—N2—H2	125.1	N3—C4—C3	105.85 (18)
C2—N3—C4	109.90 (16)	C6—C5—C4	116.3 (2)
C2—N3—H3	125.1	C6—C5—H5	121.8
C4—N3—H3	125.1	C4—C5—H5	121.8
C2—C1—H1A	109.5	C5—C6—C7	121.7 (2)
C2—C1—H1B	109.5	C5—C6—H6	119.2
H1A—C1—H1B	109.5	C7—C6—H6	119.2
C2—C1—H1C	109.5	C8—C7—C6	121.9 (2)
H1A—C1—H1C	109.5	C8—C7—H7	119.1
H1B—C1—H1C	109.5	C6—C7—H7	119.1
N2—C2—N3	108.40 (19)	C7—C8—C3	116.6 (2)
N2—C2—C1	126.37 (19)	C7—C8—H8	121.7
N3—C2—C1	125.23 (19)	C3—C8—H8	121.7
C3—N2—C2—N3	-0.6 (2)	C8—C3—C4—N3	-178.53 (17)
C3—N2—C2—C1	179.86 (19)	N2—C3—C4—N3	-0.24 (19)
C4—N3—C2—N2	0.4 (2)	N3—C4—C5—C6	178.52 (19)
C4—N3—C2—C1	179.99 (19)	C3—C4—C5—C6	0.0 (3)
C2—N2—C3—C8	178.5 (2)	C4—C5—C6—C7	0.0 (3)
C2—N2—C3—C4	0.5 (2)	C5—C6—C7—C8	-0.4 (4)
C2—N3—C4—C5	-178.8 (2)	C6—C7—C8—C3	0.7 (3)
C2—N3—C4—C3	-0.1 (2)	N2—C3—C8—C7	-178.4 (2)

C8—C3—C4—C5	0.3 (3)	C4—C3—C8—C7	-0.7 (3)
N2—C3—C4—C5	178.59 (17)		

Hydrogen-bond geometry (Å, °)

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
N2—H2...O1 ⁱ	0.86	2.03	2.855 (3)	162
N3—H3...O2 ⁱⁱ	0.86	1.93	2.775 (2)	166

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