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Key indicators

Single-crystal X-ray study T = 113 K Mean σ (C–C) = 0.003 Å R factor = 0.037 wR factor = 0.082 Data-to-parameter ratio = 9.2

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

(S,S)-2,2'-(Ethane-1,2-diyldiiminio)dibutan-1-ol dinitrate

In the title structure, $C_{10}H_{26}N_2O_2^{2+}\cdot 2NO_3^{-}$, the ethambutol molecule exists in the cationic form, having two protonated amine groups, and is charge-balanced by two nitrate anions. In the crystal structure, a two-dimensional network is formed *via* intermolecular $O-H \cdots O$ and $N-H \cdots O$ hydrogen bonds.

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Comment

The title compound, (I), is the dinitrate salt of ethambutol (Bai *et al.*, 2006), which is a key intermediate in the synthesis of ethambutol hydrochloride, a widely used chiral antituberculosis agent (Fadnavis *et al.*, 1999). A view of the asymmetric unit of (I) is shown in Fig. 1. The ethambutol molecule in (I) exists in the cationic form, having two protonated amino groups. The dication is charged balanced by two nitrate anions. The bond distances and angles in the cation and the anion are normal within experiment error (Allen *et al.*, 1987). Selected torsion angles are given in Table 1.



In the crystal structure, anions and cations are connected *via* intermolecular $O-H\cdots O$ and $N-H\cdots O$ hydrogen bonds, forming a two-dimensional network perpendicular to the *a* axis (Fig. 2 and Table 2).

Experimental

The title compound was prepared by the reaction of nitric acid (0.065 g, 65%) with ethambutol (0.1 g, 0.0003 mol). Colourless single crystals were grown by slow evaporation of a solution of (I) in methanol–water (6:1).

Crystal data	
$C_{10}H_{26}N_2O_2^{2+}\cdot 2NO_3^{-}$	Z = 2
$M_r = 330.35$	$D_x = 1.373 \text{ Mg m}^{-3}$
Monoclinic, P2 ₁	Mo $K\alpha$ radiation
a = 8.769 (3) Å	$\mu = 0.12 \text{ mm}^{-1}$
b = 9.598 (4) Å	T = 113 (2) K
c = 9.583 (4) Å	Block, colorless
$\beta = 97.829 \ (6)^{\circ}$	$0.26 \times 0.24 \times 0.20$ mm
V = 799.0 (5) Å ³	

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Data collection

Bruker SMART-1000 diffractometer φ and ω scans Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996) $T_{\rm min} = 0.964, T_{\rm max} = 0.977$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.037$ $wR(F^2) = 0.082$ S = 1.072011 reflections 218 parameters

Table 1

Selected torsion angles ($^{\circ}$).

C5-N1-C1-C3	55.3 (2)	N1-C5-C6-N2	174.91 (17)
C5-N1-C1-C2	179.53 (17)	C6-N2-C7-C9	56.3 (2)
N1-C1-C3-C4	-173.70(16)	C6-N2-C7-C8	-176.23(18)
C1-N1-C5-C6	169.86 (16)		

7494 measured reflections 2011 independent reflections

 $R_{\rm int} = 0.026$

 $\theta_{\rm max} = 27.9^\circ$

refinement

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

 $\begin{array}{l} \Delta \rho_{\rm max} = 0.20 \ {\rm e} \ {\rm \AA}^{-3} \\ \Delta \rho_{\rm min} = -0.28 \ {\rm e} \ {\rm \AA}^{-3} \end{array}$

1830 reflections with $I > 2\sigma(I)$

H atoms treated by a mixture of

 $w = 1/[\sigma^2(F_o^2) + (0.0503P)^2]$ where $P = (F_o^2 + 2F_c^2)/3$

independent and constrained

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

$D - \mathbf{H} \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots \mathbf{A}$
$N2-H2D\cdots O8^{i}$	0.85 (3)	2.13 (3)	2.939 (3)	158 (2)
$N2-H2D\cdots O6^{i}$	0.85 (3)	2.54 (3)	3.041 (2)	119.0 (19)
$N1 - H1C \cdot \cdot \cdot O3^{ii}$	0.95 (3)	1.87 (3)	2.804 (2)	168 (2)
$N1-H1C\cdots O4^{ii}$	0.95 (3)	2.53 (2)	3.073 (2)	116.8 (17)
$N2-H2C\cdots O7$	1.05 (2)	1.83 (2)	2.880 (3)	176 (2)
$N2-H2C\cdots O6$	1.05 (2)	2.44 (2)	3.068 (2)	117.2 (16)
$N1 - H1B \cdots O3^{iii}$	0.90 (2)	2.06 (3)	2.877 (2)	149 (2)
$N1 - H1B \cdots O5^{iii}$	0.90(2)	2.39 (2)	3.180 (3)	146 (2)
$O2-H2 \cdot \cdot \cdot O4$	0.76 (3)	2.07 (3)	2.818 (2)	166 (3)
$O2-H2 \cdot \cdot \cdot O5$	0.76 (3)	2.63 (3)	3.262 (2)	141 (3)
$O1-H1A\cdots O7^{iii}$	0.83 (3)	1.99 (3)	2.824 (2)	173 (3)

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

In the absence of significant anomalous dispersion effects, Freidel pairs were merged; the absolute configuration is known from that of the starting material. H atoms bonded to N and O atoms were refined, with $U_{\rm iso}(\rm H) = 1.2U_{eq}(\rm N)$ or $1.5U_{eq}(\rm O)$. H atoms bonded to C atoms were included in calculated positions, with C–H = 0.98–1.00 Å and $U_{\rm iso}(\rm H) = 1.2U_{eq}(\rm C)$ or $1.5U_{eq}(\rm methyl C)$.

Data collection: *SMART* (Bruker, 1997); cell refinement: *SAINT* (Bruker, 1997); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 1997); software used to prepare material for publication: *SHELXTL*.

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Figure 1

The asymmetric unit of (I), showing the atom-numbering scheme and 30% probability displacement ellipsoids.





Part of the crystal structure of (I), with hydrogen bonds shown as dashed lines.

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