

(E)-3,3'-(Diazene-1,2-diyI)bis(1-methyl-1,4,5,6-tetrahydropyrrolo[3,4-c]pyrazol-5-iun) dinitrate dihydrate

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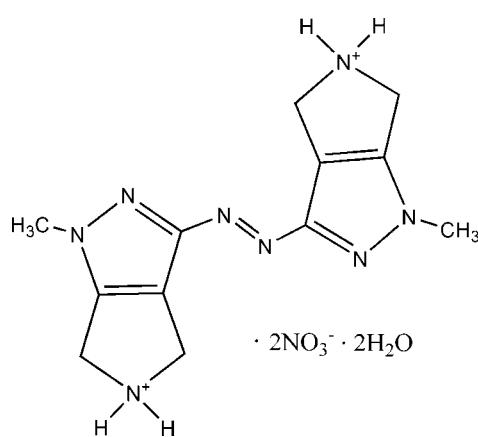
Received 28 November 2011; accepted 12 December 2011

Key indicators: single-crystal X-ray study; $T = 295\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$; R factor = 0.048; wR factor = 0.131; data-to-parameter ratio = 12.5.

The title compound, $\text{C}_{12}\text{H}_{18}\text{N}_8^{2+} \cdot 2\text{NO}_3^- \cdot 2\text{H}_2\text{O}$, was synthesized unexpectedly from 3-amino-1-methyl-1,4,5,6-tetrahydropyrrolo[3,4-c]pyrazol-5-iun chloride and cerium(IV) ammonium nitrate. The cation has a crystallographically imposed centre of symmetry. In the crystal, the ions and water molecules are linked via $\text{O}-\text{H}\cdots\text{N}$, $\text{N}-\text{H}\cdots\text{O}$ and $\text{O}-\text{H}\cdots\text{O}$ hydrogen bonds into a three-dimensional network.

Related literature

For background to potential anticancer kinase inhibitors, see: Fancelli *et al.* (2005); Gadekar *et al.* (1968). For a related structure, see: Xia *et al.* (2011).



Experimental

Crystal data

$\text{C}_{12}\text{H}_{18}\text{N}_8^{2+} \cdot 2\text{NO}_3^- \cdot 2\text{H}_2\text{O}$	$\gamma = 92.84(3)^\circ$
$M_r = 434.40$	$V = 462.64(16)\text{ \AA}^3$
Triclinic, $P\bar{1}$	$Z = 1$
$a = 6.2344(12)\text{ \AA}$	Mo $K\alpha$ radiation
$b = 7.7725(16)\text{ \AA}$	$\mu = 0.13\text{ mm}^{-1}$
$c = 9.7071(19)\text{ \AA}$	$T = 295\text{ K}$
$\alpha = 99.56(3)^\circ$	$0.22 \times 0.16 \times 0.12\text{ mm}$
$\beta = 92.49(3)^\circ$	

Data collection

Rigaku SCXmini diffractometer	1479 reflections with $I > 2\sigma(I)$
4322 measured reflections	$R_{\text{int}} = 0.034$
1811 independent reflections	

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.048$	H atoms treated by a mixture of independent and constrained refinement
$wR(F^2) = 0.131$	$\Delta\rho_{\text{max}} = 0.28\text{ e \AA}^{-3}$
$S = 1.08$	$\Delta\rho_{\text{min}} = -0.21\text{ e \AA}^{-3}$
1811 reflections	
145 parameters	

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

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
O1—H1E···N4 ⁱ	0.95 (3)	1.98 (3)	2.895 (2)	163 (2)
N2—H2B···O1 ⁱⁱ	0.90	1.94	2.802 (3)	159
N2—H2A···O1 ⁱⁱⁱ	0.90	2.44	2.970 (2)	118
N2—H2A···O3 ^{iv}	0.90	2.18	2.894 (3)	136
O1—H1F···O2	0.85 (4)	1.97 (4)	2.819 (2)	173 (3)

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

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/PC* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL/PC*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: RZ2680).

References

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supplementary materials

Acta Cryst. (2012), E68, o151 [doi:10.1107/S1600536811053347]

(E)-3,3'-(Diazene-1,2-diyl)bis(1-methyl-1,4,5,6-tetrahydropyrrolo[3,4-c]pyrazol-5-i um) dinitrate dihydrate

J.-M. Chen and H. Zhao

Comment

Tetrahydropyrrolo[3,4-c]pyrazol derivatives are used as anticancer kinase inhibitors (Xia *et al.*, 2011; Fancelli *et al.*, 2005; Gadekar *et al.*, 1968). The title compound was synthesized unexpectedly from 3-amino-1-methyl-1,4,5,6-tetrahydropyrrolo[3,4-c]pyrazol-5-i um chloride and cerium(IV) ammonium nitrate, and its crystal structure is presented herein.

The molecular structure of the title compound is shown in Fig. 1. The cation lies on a crystallographic inversion centre located at the centre of the diazene fragment. The dihedral angle between the fused pyrrole and pyrazole rings is 4.46 (12) $^{\circ}$. In the crystal structure, the ions and water molecules are linked *via* O—H \cdots N, N—H \cdots O and O—H \cdots O hydrogen bonds into a three-dimensional network. (Table 1; Fig. 2).

Experimental

3-Amino-1-methyl-1,4,5,6-tetrahydropyrrolo[3,4-c]pyrazol-5-i um chloride (0.35 g, 2 mmol) and cerium(IV) ammonium nitrate (0.28 g, 0.5 mmol) were dissolved in 95% ethanol (25 ml). The solution was filtered and left at room temperature for 10 days. Yellow prism crystals suitable for X-ray analysis were obtained by slow evaporation of the solvent.

Refinement

The water H atoms were located in a difference Fourier map and refined freely. All other H atoms were placed in calculated positions and refined using a riding model approximation, with C—H=0.96–0.92 Å, N—H=0.97 Å, and with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C}, \text{N})$ or $1.5U_{\text{eq}}(\text{C})$ for methyl groups.

Figures

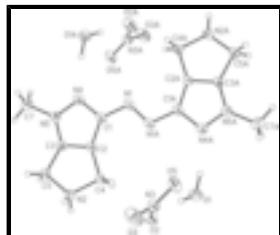


Fig. 1. The molecular structure of the title compound showing displacement ellipsoids drawn at the 30% probability level. Atoms with suffix A are generated by the symmetry operation (1-x, 1-y, 1-z).

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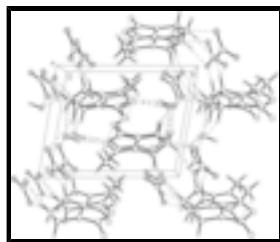


Fig. 2. Packing diagram of the title compound, showing the structure along the a axis. Dashed lines indicate hydrogen bonds.

(E)-3,3'-(Diazene-1,2-diyl)bis(1-methyl-1,4,5,6-tetrahydropyrrolo[3,4-c]pyrazol-5-ium) dinitrate dihydrate

Crystal data

$C_{12}H_{18}N_8^{2+} \cdot 2NO_3^- \cdot 2H_2O$

$M_r = 434.40$

Triclinic, $P\bar{1}$

Hall symbol: -P 1

$a = 6.2344 (12)$ Å

$b = 7.7725 (16)$ Å

$c = 9.7071 (19)$ Å

$\alpha = 99.56 (3)^\circ$

$\beta = 92.49 (3)^\circ$

$\gamma = 92.84 (3)^\circ$

$V = 462.64 (16)$ Å³

$Z = 1$

$F(000) = 228$

$D_x = 1.559$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 4309 reflections

$\theta = 3.1\text{--}27.2^\circ$

$\mu = 0.13$ mm⁻¹

$T = 295$ K

Prism, yellow

0.22 × 0.16 × 0.12 mm

Data collection

Rigaku SCXmini
diffractometer

1479 reflections with $I > 2\sigma(I)$

Radiation source: fine-focus sealed tube
graphite

$R_{\text{int}} = 0.034$

$\theta_{\max} = 26.0^\circ$, $\theta_{\min} = 3.1^\circ$

Detector resolution: 13.6612 pixels mm⁻¹

$h = -7 \rightarrow 7$

CCD_Profile_fitting scans

$k = -9 \rightarrow 9$

4322 measured reflections

$l = -11 \rightarrow 11$

1811 independent reflections

Refinement

Refinement on F^2

Primary atom site location: structure-invariant direct methods

Least-squares matrix: full

Secondary atom site location: difference Fourier map

$R[F^2 > 2\sigma(F^2)] = 0.048$

Hydrogen site location: inferred from neighbouring sites

$wR(F^2) = 0.131$

H atoms treated by a mixture of independent and constrained refinement

$S = 1.08$

$w = 1/[\sigma^2(F_o^2) + (0.060P)^2 + 0.1401P]$

1811 reflections

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

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

145 parameters $\Delta\rho_{\max} = 0.28 \text{ e } \text{\AA}^{-3}$
 0 restraints $\Delta\rho_{\min} = -0.21 \text{ e } \text{\AA}^{-3}$

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 (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.7436 (3)	0.6228 (2)	0.52005 (18)	0.0324 (4)
C2	0.8295 (3)	0.6536 (2)	0.65780 (17)	0.0301 (4)
C3	1.0187 (3)	0.7441 (2)	0.65097 (18)	0.0308 (4)
C4	0.7930 (3)	0.6264 (3)	0.80235 (18)	0.0370 (5)
H4A	0.7592	0.5043	0.8065	0.044*
H4B	0.6790	0.6959	0.8426	0.044*
C5	1.1457 (3)	0.7872 (3)	0.78510 (19)	0.0388 (5)
H5A	1.1556	0.9119	0.8199	0.047*
H5B	1.2891	0.7446	0.7779	0.047*
C7	1.2218 (3)	0.8565 (3)	0.4629 (2)	0.0467 (5)
H7A	1.3142	0.9190	0.5383	0.070*
H7B	1.1674	0.9371	0.4068	0.070*
H7C	1.3016	0.7717	0.4062	0.070*
N1	0.5569 (2)	0.5305 (2)	0.45680 (15)	0.0353 (4)
N2	1.0105 (3)	0.6894 (2)	0.87401 (16)	0.0404 (4)
H2A	0.9912	0.7595	0.9556	0.049*
H2B	1.0803	0.5968	0.8932	0.049*
N3	0.6426 (3)	0.1158 (3)	0.86872 (18)	0.0456 (5)
N4	0.8749 (3)	0.6936 (2)	0.43628 (16)	0.0363 (4)
N5	1.0436 (2)	0.7685 (2)	0.51941 (15)	0.0340 (4)
O1	0.1944 (3)	0.3662 (2)	0.86511 (17)	0.0527 (4)
O2	0.6074 (3)	0.2635 (2)	0.93400 (19)	0.0651 (5)
O3	0.8117 (3)	0.0492 (3)	0.89667 (19)	0.0733 (6)
O5	0.5152 (3)	0.0407 (3)	0.77657 (19)	0.0761 (6)
H1E	0.167 (4)	0.323 (3)	0.769 (3)	0.068 (8)*
H1F	0.323 (6)	0.344 (4)	0.886 (3)	0.084 (10)*

Atomic displacement parameters (\AA^2)

U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
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C1	0.0333 (9)	0.0308 (9)	0.0313 (9)	0.0052 (8)	-0.0060 (7)	0.0007 (7)
C2	0.0293 (9)	0.0284 (9)	0.0310 (9)	0.0027 (7)	-0.0034 (7)	0.0012 (7)
C3	0.0301 (9)	0.0286 (9)	0.0325 (9)	0.0031 (7)	-0.0016 (7)	0.0024 (7)
C4	0.0318 (9)	0.0437 (11)	0.0341 (10)	-0.0039 (8)	-0.0051 (8)	0.0065 (8)
C5	0.0311 (9)	0.0456 (11)	0.0379 (10)	-0.0057 (8)	-0.0047 (8)	0.0061 (8)
C7	0.0461 (12)	0.0463 (12)	0.0498 (12)	0.0040 (10)	0.0094 (10)	0.0122 (9)
N1	0.0359 (9)	0.0330 (8)	0.0347 (8)	0.0038 (7)	-0.0089 (6)	0.0009 (6)
N2	0.0365 (9)	0.0528 (10)	0.0302 (8)	-0.0027 (8)	-0.0063 (7)	0.0052 (7)
N3	0.0430 (10)	0.0540 (11)	0.0378 (9)	-0.0108 (9)	-0.0043 (8)	0.0078 (8)
N4	0.0403 (9)	0.0373 (9)	0.0301 (8)	0.0048 (7)	-0.0042 (7)	0.0034 (6)
N5	0.0353 (8)	0.0332 (8)	0.0336 (8)	0.0042 (7)	0.0007 (6)	0.0051 (6)
O1	0.0513 (10)	0.0694 (11)	0.0375 (8)	0.0156 (8)	0.0006 (7)	0.0056 (7)
O2	0.0619 (11)	0.0573 (11)	0.0706 (11)	0.0123 (9)	-0.0215 (8)	-0.0018 (9)
O3	0.0658 (11)	0.0726 (12)	0.0727 (12)	0.0239 (10)	-0.0173 (9)	-0.0139 (9)
O5	0.0587 (11)	0.0946 (14)	0.0639 (11)	-0.0315 (10)	-0.0129 (9)	-0.0043 (10)

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

C1—N4	1.340 (3)	C7—N5	1.451 (3)
C1—C2	1.397 (2)	C7—H7A	0.9600
C1—N1	1.398 (2)	C7—H7B	0.9600
C2—C3	1.353 (3)	C7—H7C	0.9600
C2—C4	1.479 (2)	N1—N1 ⁱ	1.258 (3)
C3—N5	1.336 (2)	N2—H2A	0.9000
C3—C5	1.474 (2)	N2—H2B	0.9000
C4—N2	1.517 (2)	N3—O5	1.221 (2)
C4—H4A	0.9700	N3—O3	1.237 (2)
C4—H4B	0.9700	N3—O2	1.250 (2)
C5—N2	1.500 (3)	N4—N5	1.344 (2)
C5—H5A	0.9700	O1—H1E	0.95 (3)
C5—H5B	0.9700	O1—H1F	0.85 (4)
N4—C1—C2	110.64 (16)	N5—C7—H7A	109.5
N4—C1—N1	116.89 (16)	N5—C7—H7B	109.5
C2—C1—N1	132.44 (17)	H7A—C7—H7B	109.5
C3—C2—C1	104.04 (16)	N5—C7—H7C	109.5
C3—C2—C4	111.36 (15)	H7A—C7—H7C	109.5
C1—C2—C4	144.60 (17)	H7B—C7—H7C	109.5
N5—C3—C2	109.44 (16)	N1 ⁱ —N1—C1	112.49 (18)
N5—C3—C5	136.16 (17)	C5—N2—C4	111.95 (14)
C2—C3—C5	114.38 (16)	C5—N2—H2A	109.2
C2—C4—N2	100.74 (14)	C4—N2—H2A	109.2
C2—C4—H4A	111.6	C5—N2—H2B	109.2
N2—C4—H4A	111.6	C4—N2—H2B	109.2
C2—C4—H4B	111.6	H2A—N2—H2B	107.9
N2—C4—H4B	111.6	O5—N3—O3	120.6 (2)
H4A—C4—H4B	109.4	O5—N3—O2	120.7 (2)
C3—C5—N2	99.89 (14)	O3—N3—O2	118.73 (18)
C3—C5—H5A	111.8	C1—N4—N5	105.67 (14)

N2—C5—H5A	111.8	C3—N5—N4	110.19 (15)
C3—C5—H5B	111.8	C3—N5—C7	128.79 (17)
N2—C5—H5B	111.8	N4—N5—C7	120.99 (16)
H5A—C5—H5B	109.5	H1E—O1—H1F	107 (3)
N4—C1—C2—C3	−0.7 (2)	N4—C1—N1—N1 ⁱ	−179.69 (18)
N1—C1—C2—C3	176.94 (18)	C2—C1—N1—N1 ⁱ	2.8 (3)
N4—C1—C2—C4	179.0 (2)	C3—C5—N2—C4	−11.9 (2)
N1—C1—C2—C4	−3.4 (4)	C2—C4—N2—C5	13.2 (2)
C1—C2—C3—N5	0.9 (2)	C2—C1—N4—N5	0.20 (19)
C4—C2—C3—N5	−178.88 (15)	N1—C1—N4—N5	−177.85 (14)
C1—C2—C3—C5	−177.86 (15)	C2—C3—N5—N4	−0.9 (2)
C4—C2—C3—C5	2.3 (2)	C5—C3—N5—N4	177.5 (2)
C3—C2—C4—N2	−9.2 (2)	C2—C3—N5—C7	−179.34 (17)
C1—C2—C4—N2	171.1 (2)	C5—C3—N5—C7	−0.9 (3)
N5—C3—C5—N2	−172.5 (2)	C1—N4—N5—C3	0.39 (19)
C2—C3—C5—N2	5.9 (2)	C1—N4—N5—C7	179.01 (16)

Symmetry codes: (i) $-x+1, -y+1, -z+1$.

Hydrogen-bond geometry (\AA , °)

$D\cdots H$	$H\cdots A$	$D\cdots A$	$D—H\cdots A$
O1—H1E···N4 ⁱ	0.95 (3)	1.98 (3)	2.895 (2)
N2—H2B···O1 ⁱⁱ	0.90	1.94	2.802 (3)
N2—H2A···O1 ⁱⁱⁱ	0.90	2.44	2.970 (2)
N2—H2A···O3 ^{iv}	0.90	2.18	2.894 (3)
O1—H1F···O2	0.85 (4)	1.97 (4)	2.819 (2)

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

supplementary materials

Fig. 1

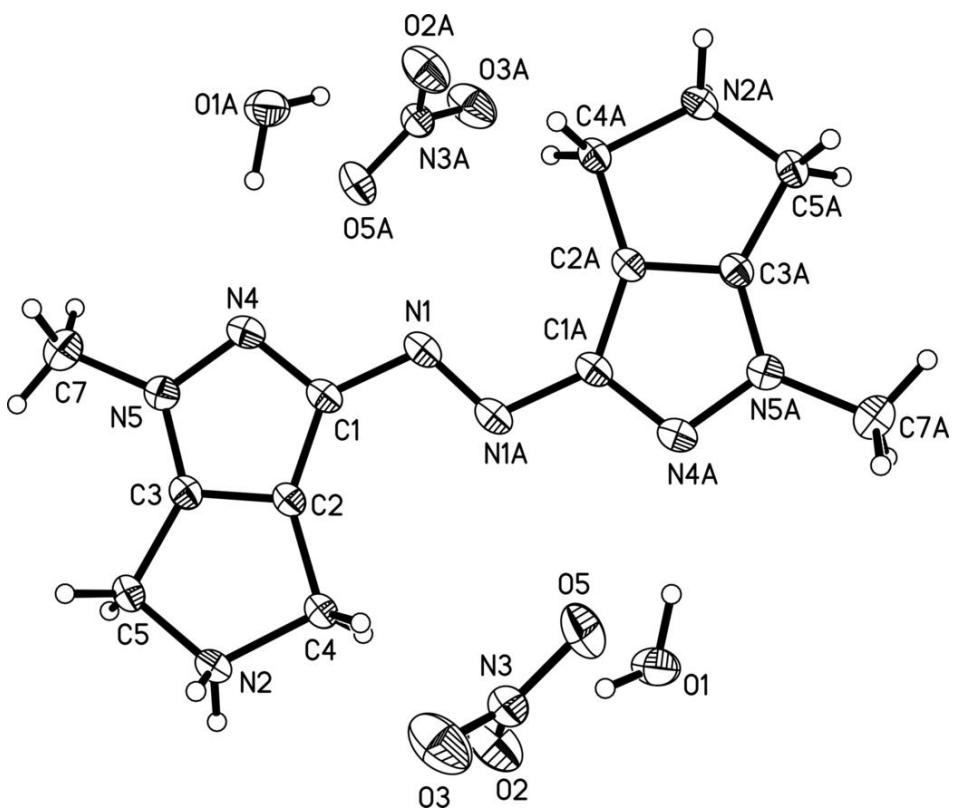


Fig. 2

