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2-Iminiumyl-1,3-diazepane-4-carboxylate

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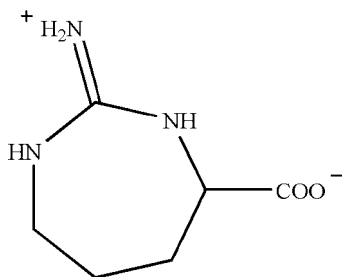
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Key indicators: single-crystal X-ray study; $T = 293$ K; mean $\sigma(\text{C}-\text{C}) = 0.005$ Å; R factor = 0.042; wR factor = 0.116; data-to-parameter ratio = 8.3.

The title compound, $\text{C}_6\text{H}_{11}\text{N}_3\text{O}_2$, is a cyclized derivative of L-arginine and the molecule is a zwitterion with the positive and negative charge residing in the guanidinium and carboxylate groups, respectively. The conformation of 1,3-diazepane ring is close to a twisted chair. One intramolecular and three intermolecular $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds stabilize the molecular conformation and the crystal structure, respectively.

Related literature

For related structures, see: Karapetyan (2008*a,b*).

Experimental

Crystal data

$\text{C}_6\text{H}_{11}\text{N}_3\text{O}_2$
 $M_r = 157.18$

Orthorhombic, $P2_12_12_1$
 $a = 6.1740$ (3) Å

$b = 8.7979$ (5) Å
 $c = 14.2036$ (7) Å
 $V = 771.51$ (7) Å³
 $Z = 4$

Mo $K\alpha$ radiation
 $\mu = 0.10$ mm⁻¹
 $T = 293$ K
 $0.23 \times 0.15 \times 0.10$ mm

Data collection

Siemens SMART CCD area-detector diffractometer
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.773$, $T_{\max} = 1.000$

3426 measured reflections
834 independent reflections
694 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.026$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.042$
 $wR(F^2) = 0.116$
 $S = 1.08$
834 reflections

100 parameters
H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.25$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.23$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N1}-\text{H1A}\cdots\text{O1}^i$	0.86	2.17	2.918 (3)	146
$\text{N3}-\text{H3A}\cdots\text{O2}^j$	0.86	2.06	2.870 (4)	157
$\text{N3}-\text{H3B}\cdots\text{O2}^{ii}$	0.86	1.95	2.788 (4)	163
$\text{N2}-\text{H2A}\cdots\text{O1}$	0.86	2.19	2.601 (3)	109

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

Data collection: SMART (Siemens, 1996); cell refinement: SMART and SAINT (Siemens, 1994); data reduction: XPREP (Siemens, 1994); program(s) used to solve structure: SHELXTL (Sheldrick, 2008); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: BX2320).

References

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Sheldrick, G. M. (1996). *SADABS*. University of Göttingen, Germany.
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supporting information

Acta Cryst. (2011). E67, o10 [https://doi.org/10.1107/S1600536810049676]

2-Iminiumyl-1,3-diazepane-4-carboxylate**Feng Yang****S1. Comment**

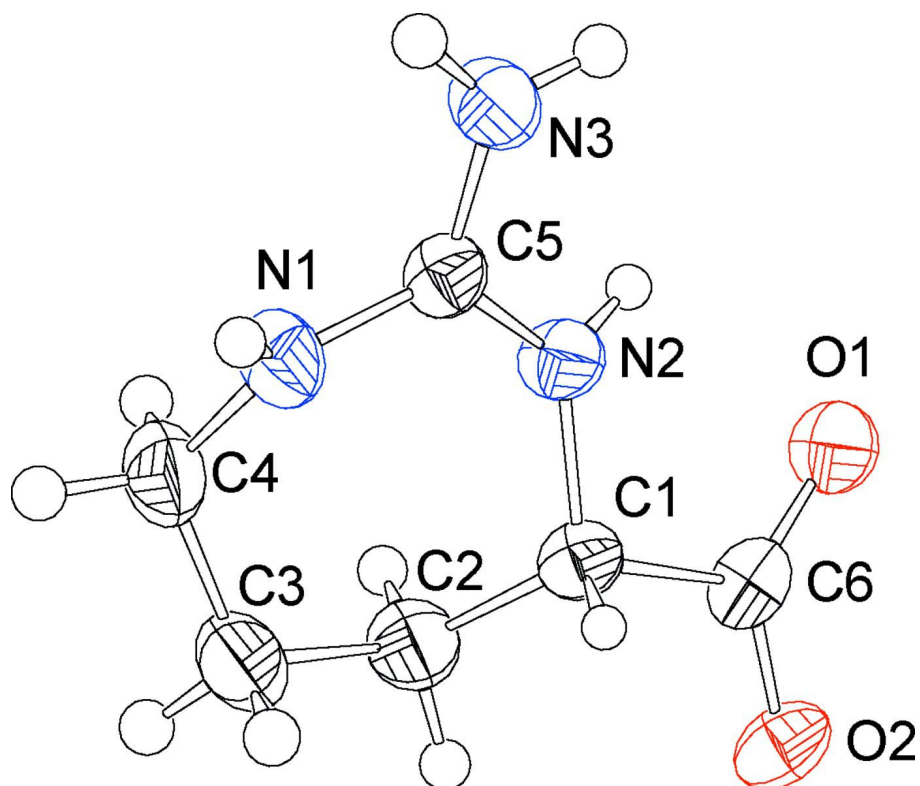
The title compound (I) was hydrothermally synthesized from *L*-Arginine *via* an unusual annulation reaction. We report here a new annulation product derived from the linear arginine molecule. The title compound is the cyclic form of *L*-Arginine and this molecule is a zwitterion with the positive and negative charge residing in the guanidinium and carboxylate groups respectively. The conformation of 1,3-diazepane ring is close to twisted chair. One intramolecular and three intermolecular N—H \cdots O hydrogen bonds stabilize the molecular conformation and the crystal structure respectively (Fig. 2). The C—N distances in the guanidinium group are obviously shorter than that of the normal C—N single bond, indicating delocalized bond of the guanidinium group.

S2. Experimental

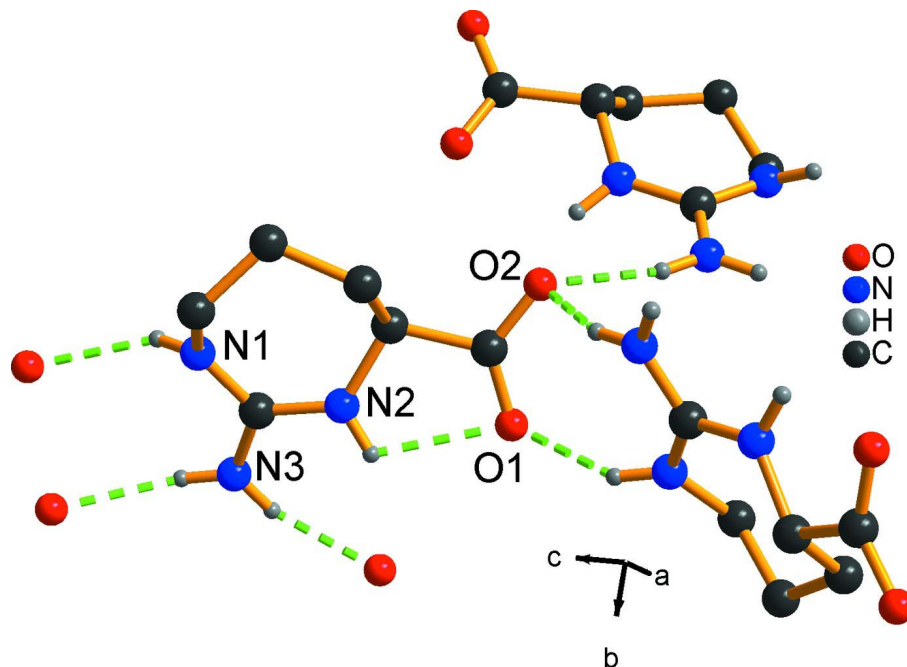
A mixture of Cu(ClO₄)₂·6H₂O (0.186 g, 0.5 mmol), *L*-Arginine (0.087 g, 0.5 mmol) and water (10 ml) was sealed in a 15 ml teflon-lined stainless steel reactor and heated to 423 K for 60 h. Colorless crystals of (I) suitable for X-ray analysis were obtained.

S3. Refinement

All H atoms were placed at calculated positions, and refined with isotropic displacement parameters, using a riding model [C—H = 0.97 Å and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$, C—H = 0.98 Å and $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C})$, N—H = 0.86 Å and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{N})$].

**Figure 1**

A view of (I) with 50% probability displacement ellipsoids; H atoms are shown as small spheres of arbitrary radii.

**Figure 2**

The hydrogen bonding interactions between the molecules. The H- atoms not involved in hydrogen bond are omitted for charity

2-iminiumyl-1,3-diazepane-4-carboxylate

Crystal data

C₆H₁₁N₃O₂ $M_r = 157.18$ Orthorhombic, $P2_12_12_1$

Hall symbol: P 2ac 2ab

 $a = 6.1740$ (3) Å $b = 8.7979$ (5) Å $c = 14.2036$ (7) Å $V = 771.51$ (7) Å³ $Z = 4$ $F(000) = 336$ $D_x = 1.353$ Mg m⁻³Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 1112 reflections

 $\theta = 2.7$ – 20.9° $\mu = 0.10$ mm⁻¹ $T = 293$ K

Prism, colorless

 $0.23 \times 0.15 \times 0.10$ mm

Data collection

Siemens SMART CCD area-detector
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

phi and ω scansAbsorption correction: multi-scan
(*SADABS*; Sheldrick, 1996) $T_{\min} = 0.773$, $T_{\max} = 1.000$

3426 measured reflections

834 independent reflections

694 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.026$ $\theta_{\max} = 25.1^\circ$, $\theta_{\min} = 2.7^\circ$ $h = -7 \rightarrow 5$ $k = -9 \rightarrow 10$ $l = -16 \rightarrow 13$

Refinement

Refinement on F^2

Least-squares matrix: full

 $R[F^2 > 2\sigma(F^2)] = 0.042$ $wR(F^2) = 0.116$ $S = 1.08$

834 reflections

100 parameters

0 restraints

Primary atom site location: structure-invariant
direct methodsSecondary atom site location: difference Fourier
mapHydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained

 $w = 1/[\sigma^2(F_o^2) + (0.0573P)^2 + 0.2708P]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\max} = 0.002$ $\Delta\rho_{\max} = 0.25$ e Å⁻³ $\Delta\rho_{\min} = -0.23$ 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	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.9643 (4)	0.4869 (3)	0.67888 (15)	0.0471 (7)
O2	0.8529 (4)	0.2509 (2)	0.64723 (15)	0.0463 (7)
N1	0.5689 (4)	0.4568 (3)	0.97693 (19)	0.0413 (8)
H1A	0.5874	0.4365	1.0356	0.050*
N2	0.7752 (5)	0.5024 (3)	0.84226 (17)	0.0383 (7)

H2A	0.8317	0.5757	0.8105	0.046*
N3	0.8387 (5)	0.6355 (3)	0.9771 (2)	0.0492 (8)
H3A	0.8106	0.6559	1.0350	0.059*
H3B	0.9404	0.6835	0.9483	0.059*
C1	0.7420 (6)	0.3583 (4)	0.7926 (2)	0.0352 (8)
H1B	0.8059	0.2765	0.8303	0.042*
C2	0.5046 (7)	0.3203 (4)	0.7748 (2)	0.0486 (10)
H2B	0.4346	0.4071	0.7457	0.058*
H2C	0.4962	0.2359	0.7310	0.058*
C3	0.3829 (7)	0.2787 (4)	0.8645 (3)	0.0558 (11)
H3C	0.2381	0.2455	0.8479	0.067*
H3D	0.4560	0.1940	0.8946	0.067*
C4	0.3667 (6)	0.4072 (4)	0.9331 (3)	0.0510 (10)
H4A	0.3033	0.4936	0.9008	0.061*
H4B	0.2671	0.3775	0.9826	0.061*
C5	0.7260 (5)	0.5309 (4)	0.9325 (2)	0.0348 (8)
C6	0.8659 (6)	0.3684 (4)	0.6983 (2)	0.0372 (8)

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0583 (16)	0.0522 (14)	0.0306 (12)	−0.0038 (14)	0.0092 (11)	0.0005 (11)
O2	0.0625 (17)	0.0467 (13)	0.0299 (11)	0.0092 (13)	0.0014 (13)	−0.0071 (10)
N1	0.0450 (18)	0.0527 (17)	0.0261 (13)	−0.0089 (15)	0.0052 (13)	0.0025 (13)
N2	0.0500 (18)	0.0391 (14)	0.0257 (13)	−0.0051 (14)	0.0053 (12)	−0.0024 (12)
N3	0.057 (2)	0.0567 (18)	0.0343 (14)	−0.0191 (17)	0.0114 (15)	−0.0135 (14)
C1	0.043 (2)	0.0356 (15)	0.0264 (16)	0.0019 (16)	−0.0015 (15)	−0.0009 (14)
C2	0.051 (2)	0.057 (2)	0.038 (2)	−0.008 (2)	0.0000 (19)	−0.0098 (17)
C3	0.049 (2)	0.061 (2)	0.057 (2)	−0.017 (2)	0.005 (2)	−0.005 (2)
C4	0.042 (2)	0.063 (2)	0.047 (2)	−0.0080 (19)	0.009 (2)	−0.0011 (18)
C5	0.0395 (19)	0.0370 (16)	0.0279 (15)	0.0017 (17)	0.0017 (15)	0.0010 (14)
C6	0.040 (2)	0.046 (2)	0.0256 (16)	0.0110 (18)	−0.0035 (16)	0.0018 (15)

Geometric parameters (Å, °)

O1—C6	1.238 (4)	C1—C2	1.524 (6)
O2—C6	1.265 (4)	C1—C6	1.545 (4)
N1—C5	1.329 (4)	C1—H1B	0.9800
N1—C4	1.462 (4)	C2—C3	1.523 (5)
N1—H1A	0.8600	C2—H2B	0.9700
N2—C5	1.340 (4)	C2—H2C	0.9700
N2—C1	1.465 (4)	C3—C4	1.496 (5)
N2—H2A	0.8600	C3—H3C	0.9700
N3—C5	1.317 (4)	C3—H3D	0.9700
N3—H3A	0.8600	C4—H4A	0.9700
N3—H3B	0.8600	C4—H4B	0.9700
C5—N1—C4	124.6 (3)	H2B—C2—H2C	107.8

C5—N1—H1A	117.7	C4—C3—C2	113.3 (3)
C4—N1—H1A	117.7	C4—C3—H3C	108.9
C5—N2—C1	126.1 (3)	C2—C3—H3C	108.9
C5—N2—H2A	116.9	C4—C3—H3D	108.9
C1—N2—H2A	116.9	C2—C3—H3D	108.9
C5—N3—H3A	120.0	H3C—C3—H3D	107.7
C5—N3—H3B	120.0	N1—C4—C3	116.5 (3)
H3A—N3—H3B	120.0	N1—C4—H4A	108.2
N2—C1—C2	113.9 (3)	C3—C4—H4A	108.2
N2—C1—C6	107.3 (3)	N1—C4—H4B	108.2
C2—C1—C6	110.2 (3)	C3—C4—H4B	108.2
N2—C1—H1B	108.4	H4A—C4—H4B	107.3
C2—C1—H1B	108.4	N3—C5—N1	120.0 (3)
C6—C1—H1B	108.4	N3—C5—N2	118.1 (3)
C3—C2—C1	112.8 (3)	N1—C5—N2	121.9 (3)
C3—C2—H2B	109.0	O1—C6—O2	126.2 (3)
C1—C2—H2B	109.0	O1—C6—C1	119.0 (3)
C3—C2—H2C	109.0	O2—C6—C1	114.8 (3)
C1—C2—H2C	109.0		

Hydrogen-bond geometry (Å, °)

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
N1—H1A \cdots O1 ⁱ	0.86	2.17	2.918 (3)	146
N3—H3A \cdots O2 ⁱ	0.86	2.06	2.870 (4)	157
N3—H3B \cdots O2 ⁱⁱ	0.86	1.95	2.788 (4)	163
N2—H2A \cdots O1	0.86	2.19	2.601 (3)	109

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