S = 4.39	Atomic scattering factors
763 reflections	from International Tables
103 parameters	for X-ray Crystallography
H atoms refined isotropically	(1974, Vol. IV, Table
$w = 1/[\sigma^2(F) + 0.0001F^2]$	2.3.1)
$(\Delta/\sigma)_{\rm max} = 0.114$	

## Table 1. Bond-length and angle comparison between pand o-nitrobenzaldehydes

ortho*	NO	N—C	C <sub>co</sub> —C <sub>aryl</sub>	C—O	C <sub>aryl</sub> -C <sub>co</sub> -O <sub>co</sub>	O-N-O
	1.226 (10)	1.468 (8)	1.490 (9)	1.200 (9)	22.2 (6)	125.0 (6)
para	1.234 (6) 1.236 (7) 1.248 (6)	1.446 (8)	1.521 (6)	1.197 (6)	116.3 (8)	120.0 (6)

\* Coppens & Schmidt (1964).

# Table 2. Fractional atomic coordinates and equivalent isotropic displacement parameters $(Å^2)$

$$U_{\rm eq} = (1/3) \sum_i \sum_j U_{ij} a_i^* a_j^* \mathbf{a}_i \cdot \mathbf{a}_j.$$

	х	y	z	$U_{eq}$
C(1)	-0.1557 (7)	0.0000	-0.8147 (4)	0.020(1)
C(2)	-0.0333 (6)	-0.1163 (15)	-0.7133 (4)	0.018 (1)
C(3)	-0.1249 (7)	-0.3210 (16)	-0.6474 (4)	0.021 (1)
C(4)	-0.3361 (7)	-0.3982 (9)	-0.6818 (4)	0.017 (1)
C(5)	-0.4572 (6)	-0.2809 (17)	-0.7881 (4)	0.023 (1)
C(6)	-0.3653 (7)	-0.0821 (15)	-0.8557 (4)	0.020(1)
N(1)	-0.0595 (6)	0.2095 (15)	-0.8852 (4)	0.027 (1)
O(2)	0.1283 (7)	0.2823 (18)	-0.8485 (4)	0.054 (1)
O(3)	-0.1645 (5)	0.3286 (15)	-0.9738 (3)	0.032 (1)
C(7)	-0.4374 (7)	-0.6224 (15)	-0.6101 (4)	0.016 (1)
O(1)	-0.3346 (5)	-0.7241 (16)	-0.5207 (3)	0.015 (1)
O(1B)	-0.6294 (19)	-0.681 (4)	-0.6482(11)	0.010 (0)

#### Table 3. Geometric parameters (Å, °)

C(1) - C(2)	1.375 (6)	C(4)—C(7)	1.521 (8)
C(1)C(6)	1.395 (6)	C(5)—C(6)	1.377 (9)
C(1) - N(1)	1.446 (7)	N(1)O(2)	1.248 (6)
C(2) - C(3)	1.389 (9)	N(1)—O(3)	1.236 (7)
C(3)-C(4)	1.382 (6)	C(7)O(1)	1.197 (6)
C(4)—C(5)	1.412 (6)	$C(7) \longrightarrow O(1B)$	1.256 (13)
C(2)C(1)C(6)	123.0 (4)	C(4)C(5)C(6)	119.4 (4)
C(2) = C(1) = N(1)	118.5 (4)	C(1)—C(6)—C(5)	118.4 (4)
C(6) - C(1) - N(1)	118.4 (4)	C(1) - N(1) - O(2)	118.3 (5)
C(1) - C(2) - C(3)	118.3 (4)	C(1) - N(1) - O(3)	121.4 (4)
C(2) - C(3) - C(4)	120.0 (4)	O(2) - N(1) - O(3)	120.0 (7)
C(3) - C(4) - C(5)	120.7 (5)	C(4)C(7)O(1)	119.1 (4)
C(3)-C(4)-C(7)	120.3 (4)	C(4) - C(7) - O(1B)	116.3 (8)
C(5) - C(4) - C(7)	118.8 (4)	O(1) - C(7) - O(1B)	124.6 (9)

Data collection: Siemens R3m/V software. Cell refinement: SHELXTL-Plus (Sheldrick, 1990). Data reduction: SHELXTL-Plus. Program(s) used to solve structure: SHELXTL-Plus. Program(s) used to refine structure: SHELXTL-Plus. Molecular graphics: SHELXTL-Plus. Software used to prepare material for publication: manual.

Lists of structure factors, anisotropic displacement parameters, Hatom coordinates and complete geometry have been deposited with the IUCr (Reference: CR1179). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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# The Linear Tripeptide L-Alanylglycyl-Lalanine

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#### Abstract

L-Alanylglycyl-L-alanine, C<sub>8</sub>H<sub>15</sub>N<sub>3</sub>O<sub>4</sub>, exists as zwitterion in the crystal with the N terminus protonated and the C terminus in an ionized form. Both the peptide units are in *trans* configurations and deviate significantly from planarity. Backbone torsion angles are  $\psi_1 = 172.7$  (2),  $\omega_1 = -178.2$  (2),  $\varphi_2 = 91.7$  (2),  $\psi_2 = -151.9$  (2),  $\omega_2$ = -176.9 (2),  $\varphi_3 = -71.3$  (2),  $\psi_{31} = -7.0$  (3) and  $\psi_{32}$ = 172.4 (2)°. The protonated NH<sub>3</sub><sup>+</sup> group forms three hydrogen bonds with atoms of symmetry-related molecules.

#### Comment

Accurate geometric parameters of simple peptides are extensively utilized in theoretical conformational studies of polypeptides and proteins. In addition, knowledge of the hydrogen-bonding scheme and van der Waals contacts in simple peptides will aid understanding of the secondary structures of polypeptides and proteins. We report here the crystal structure of L-alanylglycyl-Lalanine, AGA.



The atomic numbering and molecular conformation of the AGA molecule are shown in Fig. 1. AGA exists as a zwitterion with terminal  $NH_3^+$  and  $COO^-$  groups. Both peptide units (C1A, C1', N2, C2A and O1; C2A,

C2', N3, C3A and O2') show deviations of 0.04 Å from their respective least-squares planes. The peptide backbone assumes a folded conformation with torsion angles  $\psi_1 = 172.7(2), \varphi_2 = 91.7(2), \psi_2 = -151.9(2),$  $\varphi_3 = -71.3(2), \, \omega_1 = -178.2(2), \, \omega_2 = -176.9(2), \, \psi_{31}$ = -7.0(3) and  $\psi_{32} = 172.4(2)^{\circ}$ . The dihedral angle between the two peptide planes is  $76.7(1)^{\circ}$  and the carboxyl group makes a dihedral angle of  $73.6(2)^{\circ}$  with the adjacent peptide unit.



Fig. 1. Atomic numbering and molecular conformation of AGA. Displacement ellipsoids are drawn at the 50% probability level.

The crystal packing of AGA is shown in Fig. 2. The NH<sup>‡</sup> group forms three hydrogen bonds with the O1' and carboxyl O3' atoms of symmetry-related molecules. The packing is further stabilized by intermolecular hydrogen bonds between N2 and O3, and N3 and O3' atoms. The hydrogen-bonding geometry is given in Table 3. The conformation of AGA is different from those of AAA (Fawcett, Camerman & Camerman, 1975), which has conformational angles for an antiparallel pleated-sheet arrangement, and AGG (Subramanian & Lalitha, 1983), which has a poly(glycine)-II type conformation.



Fig. 2. Stereoview of the crystal packing of AGA viewed down the b C3A axis. Hydrogen bonds are shown by broken lines.

#### Experimental

Crystals were grown by direct evaporation from an aqueous solution of the title compound (Sigma Chemical Company, USA).

Cu  $K\alpha$  radiation

 $\mu = 0.949 \text{ mm}^{-1}$ 

T = 293 (2) K

Parallelopiped

Colourless

 $R_{\rm int} = 0.0129$  $\theta_{\rm max} = 71.91^{\circ}$ 

 $h = -5 \rightarrow 4$ 

 $k = -9 \rightarrow 17$ 

 $l = -9 \rightarrow 18$ 

2 standard reflections

frequency: 33 min

intensity variation: 4%

Cell parameters from 24 reflections

 $0.8\,\times\,0.12\,\times\,0.08$  mm

 $\lambda = 1.5418 \text{ Å}$ 

 $\theta = 12 - 8^{\circ}$ 

Crystal data

$$C_8H_{15}N_3O_4$$
  

$$M_r = 217.23$$
  
Orthorhombic  

$$P2_{12}2_12_1$$
  

$$a = 4.747 (7) \text{ Å}$$
  

$$b = 14.399 (2) \text{ Å}$$
  

$$c = 15.191 (2) \text{ Å}$$
  

$$V = 1038.5 (15) \text{ Å}^3$$
  

$$Z = 4$$
  

$$D_3 = 1.389 \text{ Mg m}^{-3}$$

Data collection

Enraf-Nonius CAD-4 diffractometer  $\omega$ -2 $\theta$  scans Absorption correction:

none 1226 measured reflections

- 1220 independent reflections
- 1141 observed reflections
  - $[I > 2\sigma(I)]$

Refinement

NI CIA C1BC1′ 01 N2 C2A C2′ 02' N3

C3B

 $\Delta \rho_{\rm max} = 0.149 \ {\rm e} \ {\rm \AA}^{-3}$ Refinement on  $F^2$  $\Delta \rho_{\rm min} = -0.134 \ {\rm e} \ {\rm \AA}^{-3}$ R(F) = 0.0279 $wR(F^2) = 0.0780$ Extinction correction: SHELXL93 (Sheldrick, S = 1.0571220 reflections 1993) Extinction coefficient: 152 parameters 0.0045 (10) All H-atom coordinates (but Atomic scattering factors not occupancies or U's) from International Tables riding on the coordinates for Crystallography (1992, of the parent atom Vol. C, Tables 4.2.6.8 and  $w = 1/[\sigma^2(F_a^2) + (0.0510P)^2]$ 6.1.1.4) + 0.1903Pwhere  $P = (F_o^2 + 2F_c^2)/3$ Absolute configuration: Flack (1983) parameter  $(\Delta/\sigma)_{\rm max} = 0.009$ -0.11(31)

Table 1. Fractional atomic coordinates and equivalent isotropic displacement parameters (Å<sup>2</sup>)

## $U_{\text{eq}} = (1/3) \sum_{i} \sum_{j} U_{ij} a_i^* a_j^* \mathbf{a}_i \cdot \mathbf{a}_j.$

x	v	ε	$U_{eq}$
1.2272 (4)	0.83935 (11)	0.62268 (10)	0.0315 (4)
1.1514 (4)	0.90207 (12)	0.54878 (11)	0.0265 (4)
1.1777 (8)	1.00243 (14)	0.5771 (2)	0.0551 (8)
1.3323 (4)	0.87763 (11)	0.46942 (10)	0.0216(3)
1.5264 (3)	0.82158 (9)	0.47548 (8)	0.0321 (3)
1.2623 (3)	().92150 (10)	().39590 (9)	0.0267 (3)
1.4061 (4)	().90477 (11)	0.31382 (11)	0.0257 (4)
1.2632 (4)	0.82911 (12)	0.26043 (11)	0.0241 (4)
1.0067 (3)	0.81594 (11)	0.26477 (11)	0.0411 (4)
1.4297 (3)	0.78090 (10)	0.20660 (9)	0.0239 (3)
1.3118 (4)	0.70997 (12)	0.14924 (11)	0.0240 (4)
1.5176 (5)	0.6855 (2)	0.07654 (14)	0.0434 (6)

C3′	1.2142 (4)	0.62256 (12)	0.19913 (13)	0.0310 (4)
O3′	1.2674 (3)	0.61527 (10)	0.28009 (9)	0.0395 (4)
03	1.0856 (5)	0.56555 (11)	0.15411 (13)	0.0641 (6)

Table 2. Selected geometric parameters (Å, °)

N1—C1A	1.485 (2)	C2'—N3	1.333 (2)
C1A—C1B	1.513(3)	C2'_O2'	1.234 (3)
CIA—CI'	1.521 (2)	N3—C3A	1.455 (2)
C1'-01'	1.229(2)	C3A—C3B	1.516 (3)
C1'N2	1.325 (2)	C3A—C3′	1.541 (2)
N2—C2A	1.442 (2)	C3'O3	1.231 (2)
C2A—C2'	1.518(2)	C3'—O3'	1.260 (2)
NI-CIA-CIB	110.3 (2)	O2'—C2'—C2A	121.5 (2)
N1 - C1A - C1'	108.76 (14)	N3—C2′—C2A	115.9 (2)
C1 <i>B</i> —C1 <i>A</i> —C1′	113.6(2)	C2'-N3-C3A	120.3 (2)
01'-C1'-N2	124.3 (2)	N3—C3A—C3B	110.6(2)
01'-C1'-C1A	121.07 (15)	N3—C3A—C3′	113.25 (14)
N2-C1'-C1A	114.6(2)	C3B—C3A—C3'	111.2 (2)
C1'-N2-C2A	122.0(2)	O3—C3′—O3′	125.9 (2)
N2-C2A-C2'	111.7 (2)	O3—C3′—C3A	114.9 (2)
O2'-C2'-N3	122.6(2)	O3'—C3'—C3A	119.2 (2)

#### Table 3. Hydrogen-bonding geometry (Å, °)

$D$ — $H \cdot \cdot \cdot A$	D—H	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	$D - H \cdot \cdot \cdot A$
N1—H1N1···O1′1	0.89	2.12	2.916 (2)	150.8
N1—H2N1···O3''	0.89	1.89	2.716(3)	154.3
N3—HN3· · · O2′ <sup>II</sup>	0.86	2.12	2.922 (4)	154.2
N1—H3N1···O3′ <sup>™</sup>	0.89	2.17	3.030 (4)	161.3
N2—HN2· · ·O3 <sup>™</sup>	0.86	1.97	2.758 (3)	151.7
Symmetry codes: (i)	$x - \frac{1}{2}, \frac{3}{2} - y,$	1-z; (ii) $1+x$	$x, y, z; (iii) \frac{1}{2} + x$	$\frac{3}{2} - y, 1 - z$
(iv) $2 - x, \frac{1}{2} + y, \frac{1}{2} - \frac{1}{2}$	- z.		2	-

Data collection: *SDP* (Enraf–Nonius, 1985). Cell refinement: *SDP*. Data reduction: local program. Program(s) used to solve structure: *SHELXS86* (Sheldrick, 1985). Program(s) used to refine structure: *SHELXL93* (Sheldrick, 1993). Molecular graphics: *INSIGHT*II (Biosym Technologies, 1995); *Xtal-GX* (Hall & du Boulay, 1995). Software used to prepare material for publication: local program.

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Lists of structure factors, anisotropic displacement parameters, Hatom coordinates and complete geometry have been deposited with the IUCr (Reference: AB1324). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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# $\gamma$ -Lactones: Methyl 11-Oxa-10-oxobicyclo-[6.3.0]undec-4-ene-9-carboxylate, Methyl 11-Oxa-10-oxobicyclo[6.3.0]undec-2-ene-9carboxylate and Methyl 9-Oxa-10-oxobicyclo[6.3.0]undecane-11-carboxylate

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#### Abstract

The structures of three  $\gamma$ -lactone derivatives, methyl 11-oxa-10-oxobicyclo[6.3.0]undec-4-ene-9-carboxylate,  $C_{12}H_{16}O_4$ , methyl 11-oxa-10-oxobicyclo[6.3.0]undec-2-ene-9-carboxylate,  $C_{12}H_{16}O_4$ , and methyl 9-oxa-10-oxobicyclo[6.3.0]undecane-11-carboxylate,  $C_{12}H_{18}O_4$ , have been solved in order to determine the relative configuration of their three asymmetric stereocentres.

#### Comment

 $\gamma$ -Lactone rings are key structural features of many perfume oils (Mori, 1989). They also exhibit important biological activities such as cytotoxic, antitumoral and bactericidal properties (Hoffmann & Rabe, 1985).

Among butyrolactone derivatives,  $\alpha$ -methylene lactones are particularly important and can be prepared by decarboxylation-methylation of  $\alpha$ -carbomethoxy- or  $\alpha$ -carboxy-substituted lactones (Petragnani, Ferraz & Silva, 1986). Such substrates can be conveniently prepared by Mn<sup>III</sup>-mediated addition of acetic acid derivatives on saturated systems (de Klein, 1986; Melikyan, 1993).

The potassium salt of monomethylmalonate is of particular interest as it allows the direct synthesis of  $\gamma$ -lactones  $\alpha$ -substituted by a carbomethoxy group which are the precursors of  $\alpha$ -methylene lactones (Lamarque, Méou & Brun, 1994). When applied to 1,3-cyclooctadiene, such an approach leads to different stereoisomers as three stereocentres are formed during the reaction. While the geometry of two of them can be determined easily by NMR spectroscopy [*cis* or *trans* junction between the  $\gamma$ -lactone ring and the cyclooctene (Lamarque, Méou & Brun, 1994)], it is very difficult to assign the relative configuration of the remaining stereocentre (i.e. the C atom bearing the