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# Kazuyuki Takahashi, Taka-aki Okamura, Hitoshi Yamamoto and Norikazu Ueyama\*

Department of Macromolecular Science, Graduate School of Science, Osaka University, Toyonaka, Osaka 560-0043, Japan

Correspondence e-mail: ueyama@chem.sci.osaka-u.ac.jp

#### Key indicators

Single-crystal X-ray study T = 200 KMean  $\sigma(C-C) = 0.003 \text{ Å}$  R factor = 0.034 wR factor = 0.088 Data-to-parameter ratio = 10.5

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e. The title novel carboxylic acid derivative,  $C_9H_{17}NO_3$ , containing the *tert*-butylcarbamoyl group, has been synthesized by the reaction of glutaric anhydride and *tert*-butyl-amine. In the crystal structure, this compound forms intermolecular hydrogen bonds between the amide and carboxyl groups. However, it does not form a hydrogen-bonded dimer involving two carboxylic acid groups.

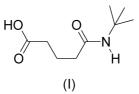
5-(tert-Butylamino)-5-oxopentanoic acid

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#### Comment

A neighbouring amide, NH, lowers the  $pK_a$  value of a carboxylic acid due to formation of an N-H···O hydrogen bond between the amide group and the anionic carboxylate O atom. This hydrogen bond prevents protonation of the carboxylate anion (Onoda *et al.*, 2003, 2004). To date, these effects have been discussed mainly with respect to acylaminobenzoic acid derivatives.

In the present study, we have synthesized a novel amidated carboxylic acid derivative, 5-(*tert*-butylamino)-5-oxopentanoic acid, (I), from glutaric anhydride and *tert*-butylamine. In the crystal structure, the alkyl backbone of (I) displays an extended conformation (Fig. 1). There is no hydrogen-bonded dimer involving the formation of intermolecular  $O-H\cdots O=C$  hydrogen bonds between two carboxylic acid groups, as discussed in an early survey (Jeffrey & Sax, 1963). This situation is rare for carboxylic acid structures.

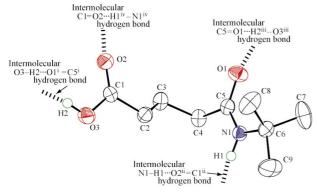


Generally, amide groups, as in nylon, interact with each other, but in (I) there are no such interactions. There are, however, intermolecular interactions between the carboxylic acid group and the amide group, *viz*. N1–H1···O2<sup>ii</sup>=C1<sup>ii</sup> [N1···O2<sup>ii</sup> 2.968 (2) Å] and O3–H2···O1<sup>i</sup>=C5<sup>i</sup> [O3···O1<sup>i</sup> 2.573 (2) Å] (Fig. 2; symmetry codes as in Table 2). Such interactions have often been reported (*e.g.* Chauhan *et al.*, 1979; Kennard *et al.*, 1982).

## **Experimental**

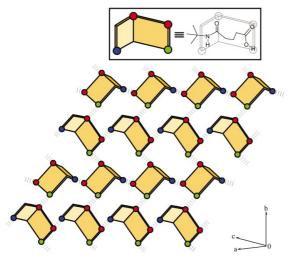
Glutaric anhydride (10.0 g, 88 mmol) in tetrahydrofuran (50 ml) was added to *tert*-butylamine (10.0 ml, 69 mmol) and the solution was stirred overnight at room temperature. After concentration of the solution, the residue was extracted with ethyl acetate. The organic layer [ethyl acetate containg (I) and an impurity] was washed with 3.5% HCl aqueous solution (30 ml), dried over anhydrous MgSO<sub>4</sub>

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### Figure 1

The structure of the title carboxylic acid, (I), with the atom-labelling scheme and with displacement ellipsoids drawn at the 50% probability level. The intermolecular hydrogen bonding is shown by dashed lines [symmetry codes: (i) and (ii) as in Table 2; (iii) 1 - x,  $y + \frac{1}{2}$ , 1 - z; (iv) x + 1, y, z].



### Figure 2

A schematic packing diagram for (I).

and concentrated, to give a white powder. This was recrystallized from ethyl acetate to give colourless block-shaped crystal of (I) (yield 43%). Spectroscopic analysis: <sup>1</sup>H NMR (d<sub>6</sub>-DMSO at 303 K,  $\delta$ , p.p.m.): 1.22 (9H, s), 1.66 (2H, m), 2.02 (2H, t), 2.17 (2H, t), 7.33 (1H, s), 12.0 (1H, s); analysis calculated for C<sub>9</sub>H<sub>17</sub>NO<sub>3</sub>: C 57.73, H 9.15, N 7.48%; found: C 57.66, H 9.08, N 7.51%.

Crystal data

$C_9H_{17}NO_3$	$D_x = 1.187 \text{ Mg m}^{-3}$
$M_r = 187.24$	Mo $K\alpha$ radiation
Monoclinic, P2 <sub>1</sub>	Cell parameters from 5001
a = 6.066 (3)  Å	reflections
b = 11.076 (6) Å	$\theta = 2.6-27.5^{\circ}$
c = 7.797 (3) Å	$\mu = 0.09 \text{ mm}^{-1}$
$\beta = 91.56 \ (4)^{\circ}$	T = 200  K
V = 523.7 (4) Å <sup>3</sup>	Block, colourless
Z = 2	$0.50$ $\times$ 0.50 $\times$ 0.20 mm
Data collection	
Rigaku RAXIS-RAPID imaging-	1128 reflections with $I > 2\sigma(I)$
plate diffractometer	$R_{\rm int} = 0.036$
$\omega$ scans	$\theta_{\rm max} = 27.5^{\circ}$
	max

plate diffractometer	$R_{\rm int} = 0.036$
$\omega$ scans	$\theta_{\rm max} = 27.5^{\circ}$
Absorption correction: none	$h = -7 \rightarrow 7$
5164 measured reflections	$k = -14 \rightarrow 14$
1246 independent reflections	$l = -10 \rightarrow 8$

#### Refinement

Refinement on $F^2$	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.034$	$w = 1/[\sigma^2(F_o^2) + (0.0647P)^2]$
$wR(F^2) = 0.088$	where $P = (F_o^2 + 2F_c^2)/3$
S = 1.03	$(\Delta/\sigma)_{\rm max} < 0.001$
1246 reflections	$\Delta \rho_{\rm max} = 0.14 \ {\rm e} \ {\rm \AA}^{-3}$
119 parameters	$\Delta \rho_{\rm min} = -0.17 \text{ e } \text{\AA}^{-3}$

### Table 1

Selected geometric parameters (Å, °).

O1-C5	1.247 (2)	N1-C5	1.323 (2)
O2-C1	1.205 (2)	N1-C6	1.484 (2)
O3-C1	1.312 (2)		
C5-N1-C6	127.37 (15)	O1-C5-C4	120.41 (16)
O2-C1-O3	123.13 (16)	N1-C5-C4	116.76 (16)
O2-C1-C2	124.01 (16)	N1-C6-C8	109.92 (16)
O3-C1-C2	112.85 (15)	N1-C6-C7	110.59 (16)
O1-C5-N1	122.78 (17)	N1-C6-C9	105.66 (16)
O2-C1-C2-C3	-35.7 (3)	C3-C4-C5-N1	110.05 (18)

#### Table 2 Hydrogen-bonding geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$O3-H2\cdots O1^{i}$	0.84	1.75	2.573 (2)	165
$N1 - H1 \cdots O2^{ii}$	0.88	2.11	2.968 (2)	163

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

H atoms were positioned geometrically and treated as riding on their parent atoms, with methyl C-H, methylene C-H, carboxylic O-H and amide N-H distances of 0.98, 0.99, 0.84 and 0.88 Å, respectively.  $U_{iso}(H)$  values were constrained to be  $1.2U_{eq}$  of the carrier atom. Rotating-group refinement was used for the methyl groups and carboxylic acid groups. In the absence of significant anomalous dispersion effects, Friedel pairs were averaged.

Data collection: PROCESS-AUTO (Rigaku Corporation, 1998); cell refinement: PROCESS-AUTO; data reduction: TEXSAN (Molecular Structure Corporation, 2000); program(s) used to solve structure: SIR92 (Altomare et al., 1994); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: TEXSAN and Mercury (Bruno et al., 2002); software used to prepare material for publication: TEXSAN and Mercury.

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