## organic papers

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#### Key indicators

Single-crystal X-ray study T = 120 KMean  $\sigma(\text{C}-\text{C}) = 0.003 \text{ Å}$  R factor = 0.039 wR factor = 0.077 Data-to-parameter ratio = 14.4

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

# $N^{\varepsilon}$ -Benzyloxycarbonyl- $N^{\alpha}$ -(2-nitrobenzenesulfonyl)-L-lysine

The title compound,  $C_{20}H_{23}N_3O_8S$ , is in an extended conformation. The COOH group does not form hydrogenbonded carboxyl dimers, but forms an intermolecular hydrogen bond with the NCOO carbonyl O atom  $[O \cdots O 2.666 (2) \text{ Å}]$ . The N atom bonded to S is pyramidal, and forms a bifurcated intramolecular hydrogen bond with carboxy and nitro O atoms. Received 22 November 2000 Accepted 28 November 2000 Online 14 December 2000

### Comment

The solid-phase peptide synthesis by the use of  $N^{\alpha}$  -protecting groups such as Fmoc (9-fluorenylmethoxycarbonyl) is widely used as described by Wenschuh *et al.* (1994). In an effort to increase the yield of the amino acid coupling reaction, the title compound, (I), was synthesized. The 2-nitrobenzenesulfonyl protecting group is smaller than the Fmoc group and is expected to increase the amino acid coupling yield due to decreased steric interactions with the nucleophile. The crystal structure determination of the title compound was carried out to confirm its successful synthesis.



The title compound is in an extended conformation. The COOH group does not form hydrogen-bonded carboxy dimers, but forms an intermolecular hydrogen bond with the carbonyl O5 atom. The lysine N1 atom is pyramidal, and forms a bifurcated intramolecular hydrogen bond with the carboxy O3 and nitro O8 atoms.

The cell dimensions of the title compound at 296 K are a = 11.8092 (13), b = 34.832 (4), c = 5.4033 (6) Å and V = 2222.6 (7) Å<sup>3</sup>.

## Experimental

The title compound was prepared by reacting N<sup> $\varepsilon$ </sup>-benzyloxycarbonyl-L-lysine with chlorotrimethylsilane (1.85 equivalents), diisopropylethylamine (2.3 equivalents) and *ortho*-nitrobenzenesulfonyl chloride (0.9 equivalents) in anhydrous dichloromethane under argon over 12 h (273 K to reflux), followed by aqueous work-up and extraction. The amino acid derivative was recrystallized by dissolving the crude material in a hot ethanol solution, followed by slow cooling to room temperature.

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#### Figure 1

The atom-numbering scheme for (I) with displacement ellipsoids at the 50% probability level.

Mo  $K\alpha$  radiation

reflections

 $\mu = 0.20 \text{ mm}^{-1}$ 

Fragment, colorless

 $0.23 \times 0.12 \times 0.10$  mm

13 981 measured reflections

4345 independent reflections

3563 reflections with  $I > 2\sigma(I)$ 

 $\theta = 2.5 - 27.5^{\circ}$ 

T = 120 K

 $R_{\rm int} = 0.050$ 

 $\theta_{\rm max} = 27.5^\circ$ 

 $l = -5 \rightarrow 5$ 

 $h = -15 \rightarrow 15$ 

 $k = -44 \rightarrow 44$ 

Cell parameters from 13981

Crystal data

 $\begin{array}{l} C_{20}H_{23}N_3O_8S\\ M_r = 465.47\\ \text{Orthorhombic, } P2_12_12\\ a = 11.7629 (2) \text{ Å}\\ b = 34.5731 (6) \text{ Å}\\ c = 5.3477 (2) \text{ Å}\\ V = 2174.8 (2) \text{ Å}^3\\ Z = 4\\ D_x = 1.422 \text{ Mg m}^{-3} \end{array}$ 

#### Data collection

KappaCCD diffractometer (with Oxford Cryosystems Cryostream cooler)  $\omega$  scans with  $\kappa$  offsets Absorption correction: multi-scan (*HKL SCALEPACK*; Otwinowski & Minor, 1997)  $T_{min} = 0.937, T_{max} = 0.980$ 

#### Refinement

Refinement on $F^2$	$w = 1/[\sigma^2(F_o^2) + (0.0239P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.039$	+ 0.7661P]
$wR(F^2) = 0.077$	where $P = (F_o^2 + 2F_c^2)/3$
S = 1.05	$(\Delta/\sigma)_{\rm max} = 0.001$
4345 reflections	$\Delta \rho_{\rm max} = 0.21 \ {\rm e} \ {\rm \AA}^{-3}$
302 parameters	$\Delta \rho_{\rm min} = -0.24 \text{ e} \text{ Å}^{-3}$
H atoms treated by a mixture of	Extinction correction: SHELXL97
independent and constrained	Extinction coefficient: 0.0039 (6)
refinement	Absolute structure: Flack (1983);
	1439 Friedel pairs
	Flack parameter = $-0.07(7)$

## Table 1

Selected geometric parameters (Å, °).

S1-O2	1.4301 (15)	O5-C7	1.238 (3)
S1-O1	1.4323 (15)	O6-C7	1.336 (3)
S1-N1	1.6065 (18)	N1-C1	1.473 (3)
S1-C8	1.791 (2)	N2-C7	1.326 (3)
O3-C2	1.207 (3)	N2-C6	1.464 (3)
O4-C2	1.331 (3)		
S1-N1-C1-C3	101.17 (18)	O8-N3-C9-C8	-23.1(3)
N1-C1-C2-O3	5.7 (3)	O6-C14-C15-C20	-44.1 (3)

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

	D II	TT 4	D 4	
$D - \mathbf{H} \cdots \mathbf{A}$	D-H	$H \cdots A$	$D \cdots A$	$D - H \cdots A$
$O4{-}H40{\cdots}O5^i$	0.93 (3)	1.75 (3)	2.666 (2)	172 (3)
$N1-H1N\cdots O3$	0.80 (3)	2.26 (3)	2.659 (3)	111 (2)
$N1 - H1N \cdots O8$	0.80 (3)	2.19 (3)	2.843 (3)	138 (2)
$N2-H2N\cdots O5^{ii}$	0.87 (2)	2.41 (3)	3.233 (3)	160 (2)
$C1-H1\cdots O1^{ii}$	1.00	2.22	3.197 (3)	167
$C11 - H11 \cdots O1^{iii}$	0.95	2.42	3.254 (4)	146

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

H atoms on C atoms were placed in calculated positions with C–H bond distances in the range 0.95–1.00 Å and  $U_{\rm iso} = 1.2U_{\rm eq}$  of the attached atom, and thereafter treated as riding. H atoms on O and N atoms were placed by difference maps and refined individually. The absolute configuration was determined by refinement of the Flack (1983) parameter, and corresponds with that of L-lysine.

Data collection: *COLLECT* (Nonius, 2000); cell refinement: *DENZO* and *SCALEPACK* (Otwinowski & Minor, 1997); data reduction: *DENZO* and *SCALEPACK* (Otwinowski & Minor, 1997); program(s) used to solve structure: direct methods using *SIR* (Altomare *et al.*, 1993); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3* (Farrugia, 1997); software used to prepare material for publication: *SHELXL97*.

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