

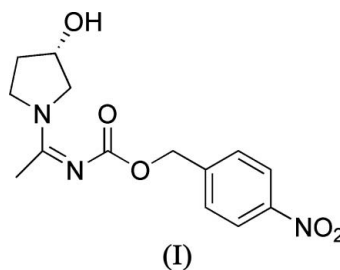
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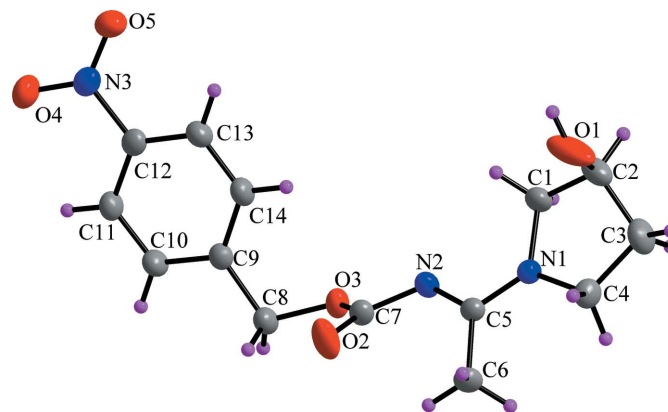
Key indicators

Single-crystal X-ray study
 $T = 293$ K
Mean $\sigma(\text{C}-\text{C}) = 0.004$ Å
 R factor = 0.039
 wR factor = 0.108
Data-to-parameter ratio = 8.4For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.**(S)-3-Hydroxy-1-(N-p-nitrobenzyloxycarbonyl-
acetimidoyl)pyrrolidine**The crystal structure of the title compound, $\text{C}_{14}\text{H}_{17}\text{N}_3\text{O}_5$, is
stabilized by intermolecular $\text{O}-\text{H}\cdots\text{N}$ hydrogen bonds.Received 23 February 2006
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Comment

Carbapenem antibiotics represent a promising set of
compounds which exhibit excellent antiviral properties
(Albers-Schonberg *et al.*, 1978; Kondo *et al.*, 1997). To date,
several carbapenem antibiotics have appeared on the market,
e.g. imipenem, panipenem, meropenem, biapenem, ertapenem
and doripenem. During our research on panipenem (Miyadera
et al., 1983), we synthesized and crystallized the title inter-
mediate, (I), and present its crystal structure here.The hydroxyl group is in an axial position of the pyrrolidine
ring. The crystal structure is stabilized by intermolecular $\text{O}-\text{H}\cdots\text{N}$
hydrogen bonds (Table 1 and Fig. 2).

Experimental

The title compound was synthesized according to the method of
Miyadera *et al.* (1995). Solvents were chromatographically pure, and
triethylamine was dried using a molecular sieve before use. 3-**Figure 1**
The molecular structure of the title compound, showing 30% probability
displacement ellipsoids (spheres of arbitrary radius for the H atoms).

Hydroxypyrrolidine hydrochloride, ethyl acetimidate hydrochloride and *p*-nitrobenzoyloxycarbonyl chloride were dried under vacuum for 4 h before use. Other chemicals were used as received without further purification. To a suspension of 3-hydroxypyrrolidine hydrochloride (1.23 g) in ethanol (10 ml) was added triethylamine (1.4 ml), followed by ethyl acetimidate hydrochloride (1.23 g). The mixture was then stirred at room temperature for 1 h. At the end of this time, the solvent was distilled off and dichloromethane (10 ml) was added. The mixture was ice-cooled, and then *p*-nitrobenzoyloxycarbonyl chloride (2.2 g) was added. Triethylamine (1.4 ml) was added dropwise and the whole mixture was stirred for 1 h. The mixture was then extracted with water, washed with brine and dried. The solvent was distilled off to obtain the title compound as a light-yellow solid (51% yield). Crystals suitable for X-ray diffraction were obtained by slow evaporation of a solution in ethyl acetate (m.p. 391–393 K).

Crystal data

$C_{14}H_{17}N_3O_5$	$Z = 2$
$M_r = 307.31$	$D_x = 1.393 \text{ Mg m}^{-3}$
Monoclinic, $P2_1$	Mo $K\alpha$ radiation
$a = 7.2123 (5) \text{ \AA}$	$\mu = 0.11 \text{ mm}^{-1}$
$b = 9.2723 (6) \text{ \AA}$	$T = 293 (2) \text{ K}$
$c = 10.9592 (7) \text{ \AA}$	Block, colourless
$\beta = 92.0580 (10)^\circ$	$0.48 \times 0.45 \times 0.42 \text{ mm}$
$V = 732.42 (8) \text{ \AA}^3$	

Data collection

Bruker SMART 1000 CCD area-detector diffractometer	4426 measured reflections
ω scans	1703 independent reflections
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)	1575 reflections with $I > 2\sigma(I)$
$T_{\min} = 0.950$, $T_{\max} = 0.956$	$R_{\text{int}} = 0.023$
	$\theta_{\text{max}} = 27.0^\circ$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0559P)^2 + 0.176P]$
$R[F^2 > 2\sigma(F^2)] = 0.039$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.108$	$(\Delta/\sigma)_{\text{max}} < 0.001$
$S = 1.05$	$\Delta\rho_{\text{max}} = 0.37 \text{ e \AA}^{-3}$
1703 reflections	$\Delta\rho_{\text{min}} = -0.29 \text{ e \AA}^{-3}$
202 parameters	Extinction correction: SHELXTL (Sheldrick, 1997)
H-atom parameters constrained	Extinction coefficient: 0.155 (13)

Table 1

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$O1-H1\cdots N2^i$	0.82	2.27	2.879 (3)	132

Symmetry code: (i) $-x, y + \frac{1}{2}, -z + 1$.

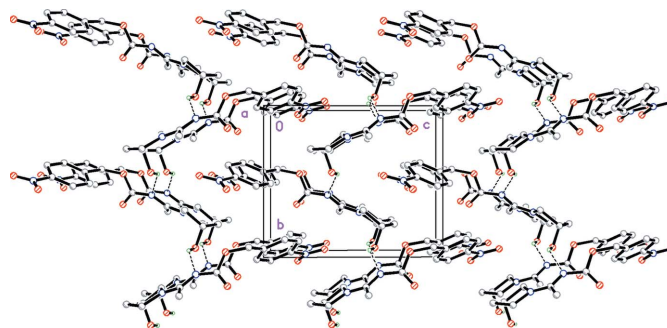


Figure 2

The crystal packing of the title compound, viewed along the *a* axis. Hydrogen bonds are shown as dashed lines. H atoms not involved in hydrogen bonding have been omitted.

All H atoms were placed in geometrically idealized positions, with $C-H = 0.96 \text{ \AA}$, and refined as riding, with $U_{\text{iso}}(H) = 1.5U_{\text{eq}}(C)$. In addition, the torsion angles about the methyl and hydroxyl groups were refined. In the absence of significant anomalous dispersion effects, Friedel pairs were averaged.

Data collection: SMART (Siemens, 1996); cell refinement: SAINT (Siemens, 1996); data reduction: SAINT; program(s) used to solve structure: SHELXTL (Sheldrick, 1997); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL.

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