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# **Structure Reports Online**

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#### **Key indicators**

Single-crystal X-ray study T = 173 KMean  $\sigma(\text{C-C}) = 0.003 \text{ Å}$  R factor = 0.049 wR factor = 0.142Data-to-parameter ratio = 17.5

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

# (2S\*,3R\*,4S\*,5R\*)-3-(S\*-1-Benzyloxyethyl)-4-methyl-4-nitro-5-phenylproline methyl ester

At 173 K, the five-membered ring of the title compound,  $C_{22}H_{26}N_2O_5$ , has an envelope conformation. The amine group is involved in both intramolecular and intermolecular hydrogen bonds, the latter linking the molecules into centrosymmetric dimers.

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#### Comment

The title compound, (3c), is one of three diastereoisomeric products, (3a-c), obtained from the [3+2]-cycloaddition of the chiral nitroalkene (1) with the silver azomethine ylide derived from imine (2) (Ayerbe *et al.*, 1998). Diastereoisomer (3a) could not be isolated; its absolute configuration was assumed by analogy with the compounds obtained in our previous work (Ayerbe *et al.*, 1998). Compound (3b) did not give crystals that were suitable for X-ray diffraction. Its relative configuration was determined by NOE experiments. The minor diastereoisomer (3c) was successfully crystallized from ethanol and fully characterized.

The puckering parameters (Cremer & Pople, 1975) for the five-membered ring are:  $q_2 = 0.463$  (2) Å and  $\varphi_2 = 321.4$  (2)°. The latter parameter is close to a value (324°) that is appropriate for an envelope conformation. The envelope flap is formed by C5, which lies 0.699 (3) Å from the plane defined by atoms N1, C2, C3 and C4. This puckering causes significant contraction of the N1—C5—C4 angle (Table 1).

The amine group of the molecule forms long, weak bifurcated hydrogen bonds with the carbonyl-O atom of the ester substituent (Table 2). One interaction is intramolecular and completes a closed five-membered loop with a graph-set motif of S(5) (Bernstein *et al.*, 1995). The second is an intermolecular interaction with a neighbouring molecule, which is related to the original molecule by a centre of inversion. The intermolecular interactions thus link the molecules into hydrogen-bonded dimers which have a graph-set motif of  $R_3^2(10)$ .

## **Experimental**

The title compound was prepared according to the procedure of Ayerbe *et al.* (1998). The reaction gave a mixture of three diaster-eoisomers, (3*a*–*c*), in the proportions 59:29:12, which were separated

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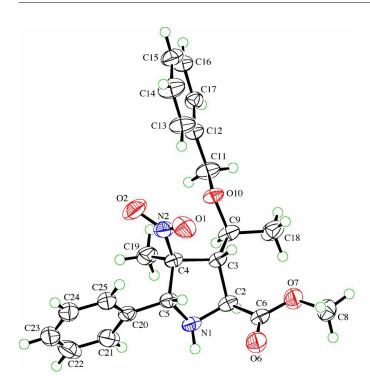


Figure 1 View of the molecule of (3c) showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 50% probability level. H atoms are represented by circles of arbitrary size.

by flash chromatography (ethyl acetate/hexane, 1:10). Suitable crystals were obtained by evaporation of an ethanol solution (m.p. 454–455 K). Spectroscopic analysis: IR (KBr,  $\nu$  cm $^{-1}$ ): 3328, 1722, 1535, 1381;  $^{1}$ H NMR (CDCl<sub>3</sub>,  $\delta$ , p.p.m.): 7.31–7.14 (m, 10 H), 4.96 ( $s_b$ , 1 H), 4.59 (d, 1 H, J = 11.5 Hz), 4.33 (d, 1 H, J = 11.5 Hz), 3.81 (s, 3 H), 3.71 (m, 3 H), 2.52 (s, 1 H), 1.30 (d, 3 H, J = 5.3 Hz), 1.26 (s, 3 H);  $^{13}$ C NMR (CDCl<sub>3</sub>,  $\delta$ , p.p.m.): 174.8, 137.6, 135.7, 128.6, 128.3, 128.2, 127.6, 127.5, 127.1, 112.3, 94.6, 73.7, 70.6, 70.1, 58.3, 57.0, 52.7, 17.0, 13.7; analysis, calculated for C<sub>22</sub>H<sub>26</sub>N<sub>2</sub>O<sub>5</sub>: C 66.30, H 6.59, N 7.03%; found: C 65.99, H 6.56, N 7.08%.

# Crystal data

Crystat auta	
$C_{22}H_{26}N_2O_5$	$D_x = 1.283 \text{ Mg m}^{-3}$
$M_r = 398.45$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/n$	Cell parameters from 25
a = 9.245 (2)  Å	reflections
b = 8.4082 (17)  Å	$\theta = 17.0  19.5^{\circ}$
c = 26.6330 (15)  Å	$\mu = 0.09 \text{ mm}^{-1}$
$\beta = 94.802 (11)^{\circ}$	T = 173 (1)  K
$V = 2063.1 (6) \text{ Å}^3$	Prism, colourless
Z = 4	$0.50 \times 0.32 \times 0.22 \text{ mm}$
Data collection	
Rigaku AFC-5R diffractometer	$\theta_{\rm max} = 27.5^{\circ}$

Rigaku AFC-5R diffractometer  $\theta_{\text{max}} = 2I.5^{\circ}$   $\omega$  scans  $h = 0 \rightarrow 12$ Absorption correction: none  $k = 0 \rightarrow 10$ 5348 measured reflections  $l = -34 \rightarrow 34$ 4719 independent reflections 3 standard reflections 2920 reflections with  $I > 2\sigma(I)$  every 150 reflections intensity decay: none

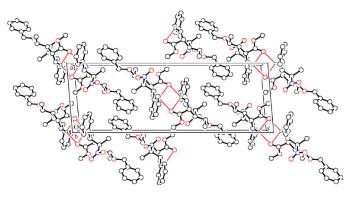


Figure 2 The molecular packing of (3c) viewed along the b axis. H atoms bonded to C atoms have been omitted for clarity. Red lines indicate the hydrogen bonding interactions.

### Refinement

refinement

-	
Refinement on $F^2$	$w = 1/[\sigma^2(F_o^2) + (0.0528P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.049$	+ 0.4415P]
$wR(F^2) = 0.142$	where $P = (F_o^2 + 2F_c^2)/3$
S = 1.06	$(\Delta/\sigma)_{\text{max}} = 0.001$
4719 reflections	$\Delta \rho_{\text{max}} = 0.27 \text{ e Å}^{-3}$
269 parameters	$\Delta \rho_{\min} = -0.19 \text{ e Å}^{-3}$
H atoms treated by a mixture of	
independent and constrained	

**Table 1** Selected geometric parameters (Å, °).

e	. ,	,	
N1-C5	1.458 (3)	C2-C3	1.563 (2)
N1-C2	1.459 (3)	C3-C4	1.549 (3)
N2-C4	1.513 (3)	C4-C5	1.550 (2)
C5-N1-C2	107.02 (15)	C3-C4-C5	101.94 (14)
N1-C2-C3	105.08 (16)	N1-C5-C4	99.45 (16)
C2-C3-C4	103.10 (15)		, ,
C5-N1-C2-C3	-29.00 (19)	C2-N1-C5-C4	46.76 (17)
N1-C2-C3-C4	-1.20(18)	C3-C4-C5-N1	-45.43(17)
C2-C3-C4-C5	28.37 (18)		` ′

 Table 2

 Hydrogen-bonding geometry ( $\mathring{A}$ ,  $^{\circ}$ ).

$D-H\cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	$D-\mathrm{H}\cdots A$
$N1-H1\cdots O6$	0.91 (2)	2.46 (2)	2.778 (2)	101.1 (16)
$N1-H1\cdots O6^{i}$	0.91 (2)	2.39 (2)	3.278 (2)	164 (2)

Symmetry code: (i) 2 - x, 1 - y, -z.

All H atoms were initially located in a difference Fourier map. The methyl-H atoms were then constrained to an ideal

geometry with C-H distances of 0.98 Å and  $U_{\rm iso}({\rm H}) = 1.5 U_{\rm eq}({\rm C})$ , but each group was allowed to rotate freely about its C-C bond. The position of the amine-H atom was refined freely along with an isotropic displacement parameter. All other H atoms were placed in geometrically idealized positions and constrained to ride on their parent atoms with C-H distances in the range 0.95–1.00 Å and  $U_{\rm iso}({\rm H}) = 1.2 U_{\rm eq}({\rm C})$ .

Data collection: MSC/AFC Diffractometer Control Software (Molecular Structure Corporation, 1991); cell refinement: MSC/AFC Diffractometer Control Software; data reduction: TEXSAN (Molecular Structure Corporation, 1999); program(s) used to solve structure: SHELXS86 (Sheldrick, 1990); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEPII (Johnson, 1976); software used to prepare material for

publication: SHELXL97 and PLATON (Spek, 2001).

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