



# Catalytic hydrogenation of vinylogous peptides: a route towards $\gamma$ -peptide foldamers

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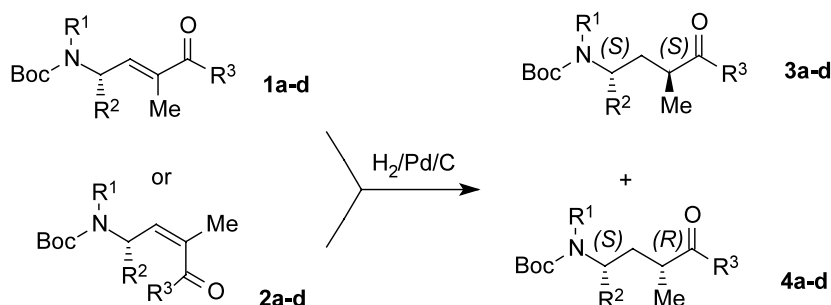
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**Abstract**—Catalytic hydrogenation over Pd/C of vinylogous aminoacids and aminoamides has been studied. The configuration of the ethylenic bond has an important effect on the diastereoselectivity. The higher selectivity is observed with the *E*-vinylogous aminoamides. The conformational preferences of the  $\alpha,\gamma$ -disubstituted  $\gamma$ -peptides have been determined. The *2S,4S*- $\gamma$ -peptide moiety induces a  $\beta$ -like folded structure stabilized by an intramolecular hydrogen bond, whereas the *2S,4R*-diastereomer assumes an open structure. © 2003 Elsevier Science Ltd. All rights reserved.

A great number of rigid molecules reproducing structural peptide units, and aiming at the design of bioactive peptide analogues with restricted flexibility, have been reported in the literature.<sup>1</sup> Most of them contain covalent rings mimicking folded regions, and more specifically the  $\beta$ -turn, a local structure spanning a tetrapeptide sequence where the carbonyl of the residue in position *i* is hydrogen-bonded to the NH in position *i*+3, thus resulting in an *N*-H $\cdots$ O=C H-bond which closes a 10-membered cycle.<sup>2</sup> Actually, these cyclic  $\beta$ -turn mimics often present different chiral centers, and their non-peptidic nature is an obstacle in the modulation of the chirality and the nature of the side chains.<sup>1c,3</sup>

We have already described the synthetic pathways to the vinylogous peptide analogues where either a *Z* or

*E*-ethylenic bond is inserted between the  $\alpha$ -carbon and the carbonyl of a peptide residue,<sup>4</sup> and reported the 3D-structure of both moieties in the solid state and in solution.<sup>4a</sup> The *E*-ethylene bond in **1d** induces an extended conformation, and the *Z*-ethylene bond in **2d** a folded structure with an N–H $\cdots$ O=C H-bond closing a nine-membered cycle. The catalytic hydrogenation of both *Z* and *E*-ethylene bonds should result in a 4-amino-2,4-disubstituted butanoic acid residue, i.e. a  $\gamma$ -peptide moiety (Scheme 1). Such a species contains two chiral centers, and we have investigated the eventual asymmetrical induction due to the chiral center in **1** and **2**, the *E* or *Z*-configuration and the C-terminal acid or amide group. We have more particularly investigated the *E* and *Z*-vinylogous derivatives listed in Table 1. Single crystals of the *Z*-vinylogous *Z*-(4*S*)-



**Scheme 1.** Catalytic hydrogenation of vinylogous peptides **1a-d** and **2a-d** into the  $\gamma$ -peptides **3a-d** and **4a-d**.

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**Table 1.** Catalytic hydrogenation of vinylogous peptides **1a–d** and **2a–d**

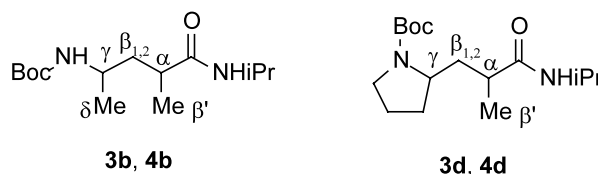
Substrates	R <sup>1</sup>	R <sup>2</sup>	R <sup>3</sup>	Conditions <sup>a</sup>	Yield (%) <sup>b</sup>	$\gamma$ -Peptide ( <i>SS/SR</i> )	<i>de</i> <sup>f</sup>
<b>1a</b>	H	Me	OH	1 bar/AcOEt–CH <sub>2</sub> Cl <sub>2</sub>	100	<b>3a/4a</b> (50/50) <sup>c</sup>	0
<b>1b</b>	H	Me	NHiPr	1 bar/AcOEt–CH <sub>2</sub> Cl <sub>2</sub>	94	<b>3b/4b</b> (77/23) <sup>d</sup>	54
<b>1b</b>	H	Me	NHiPr	50 bars/AcOEt–CH <sub>2</sub> Cl <sub>2</sub>	97	<b>3b/4b</b> (82/18) <sup>d</sup>	64
<b>2b</b>	H	Me	NHiPr	1 bar/AcOEt–CH <sub>2</sub> Cl <sub>2</sub>	87	<b>3b/4b</b> (58/42)	16
<b>1c</b>	(CH <sub>2</sub> ) <sub>3</sub>		OH	10 bars/AcOEt–CH <sub>2</sub> Cl <sub>2</sub>	98	<b>3c/4c</b> (54/46)	8
<b>1d</b>	(CH <sub>2</sub> ) <sub>3</sub>		NHiPr	50 bars/AcOEt–CH <sub>2</sub> Cl <sub>2</sub>	98	<b>3d/4d</b> (82/18) <sup>e</sup>	64
<b>2d</b>	(CH <sub>2</sub> ) <sub>3</sub>		NHiPr	50 bars/AcOEt–CH <sub>2</sub> Cl <sub>2</sub>	0	–	–
<b>2d</b>	(CH <sub>2</sub> ) <sub>3</sub>		NHiPr	10 bars/MeOH	96	<b>3d/4d</b> (78/22) <sup>e</sup>	56

<sup>a</sup> The catalytic hydrogenation was achieved in the presence of Pd on charcoal 10%.<sup>6</sup><sup>b</sup> Yield in purified products.<sup>c</sup> The absolute configuration of **3a** and **4a** was determined to be *SS* and *SR*, respectively, by comparison of the specific rotations with those reported in the literature.<sup>7</sup><sup>d</sup> The structure of **3b** was established by X-ray crystallography.<sup>e</sup> The absolute configurations were assigned using <sup>1</sup>H and <sup>13</sup>C NMR by comparing the data of pure diastereomers **3d** and **4d** with those of **3b** and **4b**.<sup>f</sup> *de* was evaluated by <sup>1</sup>H NMR (250 MHz) on the crude product.

Boc-NH-CHMe-CH=CMe-CO-NHiPr **2b** and the  $\gamma$ -peptide (2*S*,4*S*)-Boc-NH-CHMe-CH<sub>2</sub>-CHMe-CO-NHiPr **3b** were submitted to X-ray diffraction,<sup>5</sup> and the diastereomers **3b** and **4b** were studied in solution by IR and <sup>1</sup>H NMR spectroscopy.

The results of the catalytic hydrogenation are listed in Table 1.<sup>6</sup> Vinylogous aminoacids **1a** and **1c** are easily converted with high yields into the expected  $\gamma$ -amino acids **3a/4a** and **3c/4c**, and vinylogous aminoamides **1b** and **1d** into **3b/4b** and **3d/4d**. The diastereoselectivity of

the reaction actually depends on the structure of the substrate. No diastereoselection is observed with the vinylogous acids (*de*=0–8%), whereas the presence of an amide group increases the *de* values (16–64%). Moreover, the configuration of the ethylenic bond has an important effect on the stereoselectivity. The higher selectivity is observed with the *E*-vinylogous aminoamides **1b** and **1d** (*de*=64%). This can be related with the conformational preferences of the vinylogous aminoamides. The *E*-isomers adopt an extended conformation where the *Si*-side is sterically hindered by the

**Table 2.** <sup>1</sup>H NMR data for the diastereomeric molecules **3b–d** and **4b–d**

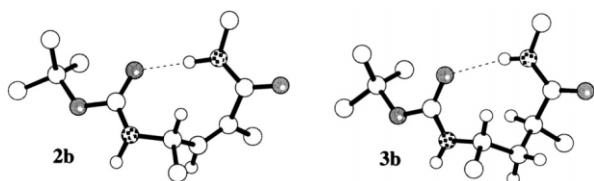
Chemical shifts (ppm)	BocNH	$\gamma$	$\delta$	$\beta_1$	$\beta_2$	$\alpha$	$\beta'$	NHiPr	$\Delta\delta$ BocNH	$\Delta\delta$ NHiPr
<b>3b</b> (CDCl <sub>3</sub> )	4.36	3.71	1.11	1.24	1.80	2.23	1.11	6.77	2.42 <sup>a</sup>	0.77 <sup>a</sup>
<b>3b</b> (C <sub>6</sub> D <sub>6</sub> )	4.20	4.06	0.90	1.09	2.02	2.43	1.42	6.97		0.57 <sup>b</sup>
<b>3b</b> (DMSO- <i>d</i> <sub>6</sub> )	6.78	3.42	0.97	1.32	1.54	2.24	0.95	7.54		
<b>4b</b> (CDCl <sub>3</sub> )	4.39	3.61	1.14	1.37	1.90	2.23	1.11	6.69	2.07 <sup>a</sup>	0.87 <sup>a</sup>
<b>4b</b> (C <sub>6</sub> D <sub>6</sub> )	4.67	3.92	1.15	1.34	2.13	2.13	1.28	5.32		2.24 <sup>b</sup>
<b>4b</b> (DMSO- <i>d</i> <sub>6</sub> )	6.46	3.41	1.02	1.26	1.61	2.22	0.96	7.56		
<b>3d</b> (CDCl <sub>3</sub> )		3.92		1.27	1.57	2.30	1.07	7.47		
<b>3d</b> (C <sub>6</sub> D <sub>6</sub> )		4.19		1.05	1.82	2.63	1.27	7.77		
<b>3d</b> (DMSO- <i>d</i> <sub>6</sub> )		3.62		1.32	1.57	2.15	0.97	7.62		
Coupling constants (Hz)	BocNH/H $\gamma$	$\gamma/\delta$	$\gamma/\beta_1$	$\gamma/\beta_2$	$\beta_1/\beta_2$	$\alpha/\beta_1$	$\alpha/\beta_2$	$\beta'/\alpha$		
<b>3b</b> (C <sub>6</sub> D <sub>6</sub> )	9.1	6.5	10.6	2.5	–13.9	2.4	11.3	6.9		
<b>3b</b> (DMSO- <i>d</i> <sub>6</sub> )	7.9	6.5	9.2	5.7	–13.4	5.9	8.9	6.9		
<b>4b</b> (C <sub>6</sub> D <sub>6</sub> )	6.4	6.5	<sup>c</sup>	<sup>c</sup>	<sup>c</sup>	<sup>c</sup>	<sup>c</sup>	6.6		
<b>4b</b> (DMSO- <i>d</i> <sub>6</sub> )	8.3	7.2	6.6	7.8	–13.3	7.3	7.0	6.8		

<sup>a</sup> Shift of the NH proton resonances when going from CDCl<sub>3</sub> to DMSO-*d*<sub>6</sub>.<sup>b</sup> Shift of the NH proton resonances when going from C<sub>6</sub>D<sub>6</sub> to DMSO-*d*<sub>6</sub>.<sup>c</sup> The medium coupling constants could not be determined exactly, because of signal broadening. The absence of large couplings denotes a flexible molecule.

Boc group (the dihydrogen approaches predominantly by the unhindered *Re*-side), whereas the intramolecular hydrogen bond in the *Z*-isomers **2b** and **2d** results in a nearly planar conformation with 2 equiv. sides for the double bond. This conformational structure agrees with the decreasing stereoselectivity and a less efficient reduction. With **2d**, the hydrogenation failed in AcOEt–CH<sub>2</sub>Cl<sub>2</sub>. The use of a polar solvent such as MeOH inhibited the formation of the intramolecular interaction and promoted the reduction and consequently **3d/4d** was obtained with an excellent yield (96%) and an interesting diastereoselectivity (*de* = 56).

It is interesting to compare the present experimental data with those from Koskinen for the hydrogenation of the vinylogous aminoester Boc-NH-CHMe-CH=CMe-COOMe.<sup>8</sup> The stereoselectivity decreases according to the sequence: R<sup>3</sup> = NHiPr (*de* = 64%) > OMe (*de* = 33%)<sup>8</sup> > OH (*de* = 0%). The control of diastereoselectivity with the *E*-vinylogous derivatives can be rationalized by the concept of allylic 1,3-strain.<sup>7</sup> The hydrogenation occurring from the *Re*-side, and leading to the (*SS*)-isomer, is clearly the result of the addition of H<sub>2</sub> onto the less hindered side in a conformation of the transition state where H $\gamma$  on the stereocenter and CH<sub>3</sub> $\beta'$  (Table 2) are eclipsed. The presence of a R<sup>3</sup> bulky group like NHiPr logically impedes the interconversion of the different conformers, and diastereoselection increases consequently (Scheme 1).

In the solid state, molecule **2b** assumes the same folded conformation as **2d**,<sup>4a</sup> and exhibits an intramolecular H-bond closing a nine-membered cycle, thus confirming the great propensity of *Z*-vinylogous residues to induce  $\beta$ -like folding (Fig. 1). More surprisingly, **3b** also adopts a folded structure where the carbons in Me-CH-CH<sub>2</sub>-CH-Me define a nearly *trans-trans* planar fragment (Fig. 1). This folded structure is retained in CH<sub>2</sub>Cl<sub>2</sub> and CHCl<sub>3</sub> on the basis of: (i) the presence of two NH stretchings at a low frequency (3320 cm<sup>-1</sup> typical of an H-bonded NH) and at a high frequency (3433 cm<sup>-1</sup> typical of a free NH); (ii) the low CO stretching frequency (1701 cm<sup>-1</sup> denoting an H-bonded Boc-CO) and (iii) the small shift of the NHiPr proton resonance (0.77 ppm from CDCl<sub>3</sub> to DMSO-*d*<sub>6</sub>) translating the small solvent accessibility of the NHiPr group (Table 2). In C<sub>6</sub>D<sub>6</sub>, a solvent which is known to spread about the proton resonances due to current ring effect, the two sets of high and small vicinal coupling constants (Table 2) confirm the *trans-trans* conformation



**Figure 1.** Crystal molecular structures of the *Z*-vinylogous peptide **2b** and the  $\gamma$ -peptide **3b**. The methyl hydrogens and the disordered isopropyl methyl carbons have been omitted for clarity.

of the Me-CH-CH<sub>2</sub>-CH-Me fragment. Under the same experimental conditions, diastereomer **4b** exhibits completely different data (high frequencies of the NH stretchings at 3433 cm<sup>-1</sup> and Boc-CO stretching at 1706 cm<sup>-1</sup>, and the large shift of 2.24 ppm for the NHiPr proton resonance) that indicate an open extended conformation. In contrast with the *Z*-vinylogous analogue **2b** which at least partly retains a folded conformation in DMSO-*d*<sub>6</sub>,<sup>4a</sup> both **3b** and **4b** only exhibit a very broad NH stretching absorption about 3250 cm<sup>-1</sup>, indicating that DMSO solvation of the NHs is favored over the aforementioned NH $\cdots$ O=C H-bond. The evolution of the vicinal coupling constants (decreasing of the high values and increasing of the small values) confirms the flexibility of the molecular backbone in this strong solvating medium.<sup>4a</sup>

The stereoselectivity of catalytic hydrogenation of the ethylenic bond in vinylogous peptides depends both on the *Z* or *E*-configuration of the ethylene bond and the nature of the contiguous acid or ester<sup>8</sup> or amide group. In most cases, the stereoselectivity (*de* = 0–64%) favors the (2*S*,4*S*)-stereomer, and the best results are obtained with the *E*-ethylene bond contiguous to an amide bond. Both *Z*-vinylogous and (2*S*,4*S*)-reduced vinylogous (2*S*,4*S*- $\gamma$ -peptide) motifs in a peptide chain induce a  $\beta$ -like folded structure stabilized by an intramolecular hydrogen bond. The stability of this turn greatly depends on the chirality since the (2*S*,4*R*)-diastereomer assumes an open structure. Recently, the mirror image of the above folded structure has been observed for (2*R*,3*R*,4*R*,5*R*,6*R*,7*R*,8*R*)-Boc-NH-CHMe-CHMe-CHMe-CO-NH-CHMe-CHMe-CHMe-CHiPr-COOBzl, thus confirming the folding tendency of  $\gamma$ -peptides having the same absolute chirality in both positions 2 and 4.<sup>9</sup>

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5. Crystal data: **2b**: tetragonal;  $P4_12_12$ ;  $a=b=10.075(2)$  Å,  $c=35.226(9)$  Å;  $Z=8$ ,  $d_{\text{calcd}}=0.978$  g cm<sup>-3</sup>; 1990 reflections;  $R=0.0703$  (each isopropyl methyl occupies three disordered positions) CCDC 200217. The main torsional angles: C–O–CO–NH  $-176^\circ$ , O–CO–NH–CH  $172^\circ$ , CO–NH–CH–CH  $-76^\circ$ , NH–CH–CH=C  $122^\circ$ , CH–CH=C–CO  $0^\circ$ , CH=C–CO–NH  $-57^\circ$ , C–CO–NH–CH  $-178^\circ$ . **3b**: tetragonal;  $P4_12_12$ ;  $a=b=9.851(1)$  Å,  $c=36.762(5)$  Å;  $Z=8$ ,  $d_{\text{calcd}}=0.992$  g cm<sup>-3</sup>; 1261 reflections;  $R=0.0679$  (each isopropyl methyl occupies two disordered positions) CCDC 200216. The main torsional angles: C–O–CO–NH  $-172^\circ$ , O–CO–NH–CH  $-173^\circ$ , CO–NH–CH–CH<sub>2</sub>  $-102^\circ$ , NH–CH–CH<sub>2</sub>–CH  $68^\circ$ , CH–CH<sub>2</sub>–CH–CO  $71^\circ$ , CH<sub>2</sub>–CH–CO–NH  $-98^\circ$ , CH–CO–NH–CH  $179^\circ$ . These data can be obtained free of charge via <http://www.ccdc.cam.ac.uk/conts/retrieving.html> (or from the Cambridge Data Centre, 12, Union Road, Cambridge CB21EZ, UK; fax: (+44)1223-336-033; or deposit@ccdc.cam.ac.uk).
6. The catalytic hydrogenation was achieved in the presence of Pd on charcoal 10% for 3 days at 20°C. The resulting  $\gamma$ -peptides **3a–d** and **4a–d** were obtained after separation of the catalyst on Celite and chromatographic purification on a silica gel column. A graduated elution gave pure diastereomers **3** and **4**. The separation of the aminoamide diastereomers **3b/4b** and **3d/4d** was easier than for aminoacids **3a/4a** and **3c/4c**. The diastereoselectivity of hydrogenation was determined by <sup>1</sup>H NMR.
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