

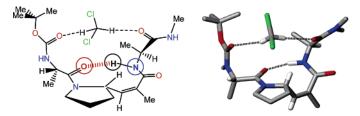
Structural Investigation of "cis" and "trans" Vinylogous Peptides: cis-Vinylog Turn in Folded cis-Vinylogous Peptides, an Excellent Mimic of the Natural β -Turn

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Various sequences of modified peptides including those containing a cis- or trans-vinylogous residue have been studied using X-ray diffraction in the solid state and ¹H NMR and IR spectroscopy in solution. A cis-vinylogous residue promotes an NH to CO intramolecular H-bond, closing a ninemembered pseudocycle that stabilizes a folded moiety that we proposed to name the cis-vinylogous turn. A trans-vinylogous residue involves an extended conformation. Two consecutive vinylogous residues retain their own structural propensity: "Xaatr"-"Xaacis" or "Xaacis"-"Xaatr" sequence is singly folded, whereas "Xaacis"-"Xaacis" sequence is doubly folded. Oligo vinylogs with all-trans or all-cis or alternating cis-trans motifs could constitute new classes of foldamers.

Introduction

The determination of the conformational features of peptides is a crucial precondition for the rational design of new bioactive peptides. Different investigations have shown that peptides that adopt specific compact conformations were particularly suited for peptide synthesis targeted toward increasing receptor selectivity, metabolic stability, and the development of potent agonists, antagonists, or enzymatic inhibitors. The most straightforward approach to induce structural restrictions is the incorporation of unnatural amino acids that adopt welldefined conformations.² Among numerous and interesting examples are various homologous amino acids such β -amino acids,³ γ-amino acids,^{3d,4} and δ-amino acids.⁵ β-Amino acid analogues such as hydrazinoacids,6 aminoxyamino acids,7 or sulfonamides8 have been also examined. In this respect, β -peptides have been extensively investigated over the past few years. They can adopt helix, sheet, or reverse turn conformations; these structures are stable, and they are adapted with short chain lengths.

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γ-Peptides have also been studied, and various secondary structures have been identified depending on degrees of substitution and configurations of γ -amino acid units. Different studies have revealed that γ^{2-4} -peptides can also adopt helical conformations in solution. 4c The conformational preferences of the $\gamma^{2,4}$ -peptides have been also determined. The 2S,4S- γ -dipeptide moiety induces a β -like folded structure by an intramolecular hydrogen bond, whereas the 2S,4R-diastereomer assumes an open structure. 4e Oligomers consisting of α,β -unsaturated γ -amino acids, so-called vinylogous peptides, have been little explored. The first example of a trans-ethylenic bond was proposed by Schreiber and co-workers as a rigid extension of a peptide residue.9 We have reported in a short communication that the insertion of a cis-properly CH= CMe group between the α-carbon and the carboxyl group of proline can be used to induce the formation of a very stable intramolecular hydrogen bond closing a ninemembered pseudocycle that involved an excellent mimic of the natural $\beta\text{-turn.}^{10}$ Recently, computational studies have shown that the introduction of a trans double bond into the backbone of the γ -amino acid constituents, which leads to vinylogous glycine, supports the folding into helices with larger H-bonded pseudocycles in the resulting peptides. 11 Synthesis and conformational studies of peptides with cis-amide bonds in trans-vinylogous proline-containing peptides have revealed the formation of 12-membered intramolecularly hydrogen-bonded structures very similar to type β -turn. ¹² Therefore, vinylogous peptides are particularly attractive structures for the construction of new peptide materials and can be used to control the structure of a peptide. A recent publication by Baldauf, Gunther, and Hofmann in this journal¹³ prompted us to publish our structural investigations concerning cis- and trans-vinylogous peptides.

In this work, we propose a detailed study of the structural features of various vinylogous peptides. The influence of the side chain, the configuration of the double bond, the position of the vinylogous amino acid in the peptide backbone, the number of vinylogous amino acid residues, and the influence of solvation are described.

TABLE 1. Synthesized Vinylogous Peptides with Their Abbreviated Codes Where "Xaa" Represents the Vinylogous Residue of the Xaa Aminoacid

"Xaa" residue	"Ala"	"Pro"	"Val"	"Gly"	"β-Ala"
Vinylogou	s Amir	oamid	es		
Boc-"Xaa ^{tr} "-NH- <i>i</i> -Pr	1a	1c	1e	1g	1i
Boc-"Xaa ^{cis} "-NH- <i>i-</i> Pr	1b	1d	1f	1ĥ	1j
Piv-"Xaa ^{tr} "-NH- <i>i-</i> Pr	2a	2c			
Piv-"Xaa ^{cis} "-NH- <i>i</i> -Pr	2b	2d			
Monovinylo	gous I	Dipeptio	des		
Boc-Pro-"Xaa ^{tr} "-NH- <i>i</i> -Pr	3a				
Boc-Pro-"Xaa ^{cis} "-NH- <i>i-</i> Pr	3b				
Piv-Pro-"Xaa ^{tr} "-NH- <i>i</i> -Pr	4a				
Boc-"Xaa ^{tr} "-Ala-NH- <i>i</i> -Pr		5c			
Boc-"Xaa ^{cis} "-Ala-NH- <i>i-</i> Pr		5d			
Boc-"Xaa ^{tr} "-Ala-NH-Me		6c			
Boc-"Xaa ^{cis} "-Ala-NH-Me		6d			
Monovinylo	gous T	ripepti	des		
Boc-Ala-"Xaa ^{tr} "-Ala-NH- <i>i</i> -Pr		7c			
Boc-Ala-"Xaa ^{cis} "-Ala-NH- <i>i</i> -Pr		7d			
Boc-Ala-"Xaa ^{tr} "-Ala-NH-Me		8c			
Boc-Ala-"Xaa ^{cis} "-Ala-NH-Me		8d			
Divinylog	ous Di	peptide	es		
Boc-"Xaa ^{tr} "-"Ala ^{tr} "-NH- <i>i</i> -Pr		9c			
Boc-"Xaa ^{tr} "-"Ala ^{cis} "-NH- <i>i</i> -Pr		9d			
Boc-"Xaa ^{cis} "-"Ala ^{tr} "-NH- <i>i</i> -Pr		9e			
Boc-"Xaa ^{cis} "-"Ala ^{cis} "-NH- <i>i</i> -Pr	9f	9g			
Piv-"Xaa ^{tr} "-"Ala ^{tr} "-NH- <i>i</i> -Pr		10c			
Piv-"Xaa ^{cis} "-"Ala ^{tr} "-NH- <i>i</i> -Pr		10e			

In this article, for clarity of the nomenclature, we propose to note "Xaacis" and "Xaatr" the cis-(Z)- and trans-(E)-vinylogous analogues of the α -amino acid Xaa, with insertion of the cis- or trans-propenyl CH=CMe group between the α -carbon and the carbonyl.

Results

Synthesis. Different vinylogous aminoamides and vinylogous peptides (cis-monovinylogs and trans-monovinylogs), β -alanine vinylogs, and divinylog dipeptides have been synthesized. They are shown in Table 1, which shows the various synthesized vinylogous peptides together with their codes.

Recently, we demonstrated the utility of Horner reagents as starting materials for the stereoselective synthesis of *trans*- and *cis*-vinylogous amino acids. ¹⁴ In the present work, we present new developments of this methodology, especially the extension to *trans*- and *cis*-vinylogous peptides and divinylogous dipeptides. The strategy is based on stereoselective Horner reactions applied to α -aminoaldehydes with the stereochemical control of the olefination determined by the choice of the Horner reagent (Scheme 1).

trans-Vinylogous amino acids were obtained by reaction between the lithiated dianion derived from 2-diethylphosphonopropanoic acid and an aminoaldehyde Boc- X_{aa} -H. The preparation of cis-vinylogous amino acids needed another stereoselective synthesis, based on the use of two systems, either KH/ethyl 2-bis(trifluoroethyl)-phosphonopropanoate or BuLi/ethyl 2-diethylphosphonopropanoate. On the basis of our previous works, a high cis stereoselectivity was obtained with the second system when aminoaldehydes were slightly hindered (Boc-Ala-

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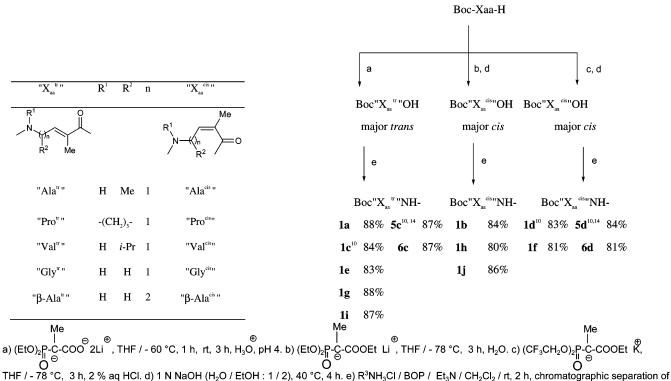
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SCHEME 1. Representative Syntheses for Vinylogous Aminoamides and Vinylogous Peptides



Z / E stereomers.

H, Boc-Gly-H, Boc- β -Ala-H), while bulkier aminoaldehydes (Boc-Pro-H, Boc-Val-H) needed Still reagent. The basic hydrolysis of pure cis-vinylogous aminoesters was effected with 1 N solution of NaOH (2.5 equiv) in H₂O/EtOH (1/2) for 2 h at 20 °C. Due to the steric hindrance of the pyrrolidine cycle, the time (4 h) and the heating (40 °C) required for completion of the reaction with Boc-"Pro"-OEt were significantly higher than those under the standard conditions.

The mixture of *cis*- and *trans*-vinylogous amino acids was coupled with the amino partners in the presence of BOP as coupling reagent and led to good yields of the desired peptides 1a-1j and 5c-6d. The reaction occurred at room temperature with 2 equiv of triethylamine in dichloromethane and was complete after being stirred for 2 h. These conditions could be applied to each amino partner (isopropylamine, H-Ala-NH-*i*-Pr, H-Ala-NH-Me). The pure trans and cis isomers could be easily obtained after a chromatographic separation.

The Boc group was removed by treatment with anhydrous HCl in Et_2O , and the amino group of the unsaturated aminoamides was acylated with pivaloyl chloride to give 2a-2d or coupled with a carboxylic partner to give 3a,3b. The best activation method was attained with BOP as coupling reagent in CH_2Cl_2 .

The vinylogous tripeptides $\mathbf{7c-8d}$ may be obtained using the same sequence: removal of the N-Boc group of $\mathbf{5c-6d}$ with HCl/coupling with a carboxylic partner using BOP as coupling reagent. The unsaturated tripeptides were easily purified on a silica gel chromatography column.

The divinylogous dipeptides 9c-10e were accessible starting from the corresponding vinylogous amino acids. The presence of the double bonds did not need modification of coupling conditions (BOP/CH₂Cl₂/Et₃N/20 °C/2h). The divinylog compounds 9c-9g were purified on silica gel without decomposition and were obtained in good yields. This purification was easy if the stereomer ratio of the ethylenic precursors was higher. The replacement of the Boc group by the pivaloyl moiety was achieved with the system HCl/Et₂O/t-BuCOCl/DIPEA and gave 10c and 10e. All the yields were high, and the reactions were relatively insensitive to the presence of the vinylogous amino acid residue. In all cases, the configuration integrity of the double bond and chiral carbons was maintained.

X-ray Diffraction. Single crystals of Piv-"Alacis"-NH-i-Pr **2b** and Boc-Ala-"Procis"-Ala-NH-Me **8d** were grown by slow evaporation of ethyl acetate/dichloromethane solution. The X-ray diffraction data set of the former was collected at room temperature on a diffractometer using Cu K α radiation from a rotating anode generator. A crystal of compound **8d** was measured at room temperature on a system equipped with a Mo sealed tube ($\lambda = 0.7107$ Å). The X-ray data were processed with WinGx¹⁶ and with HKL 2000 suite¹⁷ for the former and the latter, respectively. The crystal structures were solved by direct methods using SIR92, ¹⁸ and the models were refined by full-matrix least-squares procedures on

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 F^2 using SHELXL97. 19 The positions of H atoms attached to N atoms were located from difference maps, and the N-H bond distance was restrained to 1.03(1) Å.20 The CH hydrogen atoms were placed at a calculated position and refined using a riding model. The N-terminal tert-butyl group of Piv-"Alacis"-NH-i-Pr 2b was found to be disordered. It was modeled over two sites A and B with occupancy factors 0.769(8) and 0.231(8), respectively. In site A the three carbon atoms were refined anisotropically; in site B they were refined isotropically. In both sites the C-C bond distances were restrained to 1.51(1) A. The crystal structure determination of 8d revealed an asymmetric unit containing one Boc-Ala-"Procis"-Ala-NH-Me molecule, one dichloromethane molecule, and one disordered water molecule. This disordered water molecule was refined isotropically with a fixed occupancy factor of 0.3, and no hydrogen atoms were modeled. The main crystallographic data are as following: **2b**, $C_{14}H_{26}N_2O_2$, M = 254.37, orthorhombic, $P2_12_12$, a = 14.747(2) Å, b = 18.745(2) Å, c = 5.702(1) Å, Z = 4, $D_{\rm calcd} = 1.072 \, {\rm Mg \cdot m^{-3}}$, 1569 unique reflections, 1533 with $I > 2\sigma(I), R_1 = 0.0443 [I > 2\sigma(I)] \text{ and } wR_2 = 0.1228 [I > 0.1228]$ $2\sigma(I)$], 181 parameters; **8d**, $C_{20}H_{34}N_4O_5$, CH_2Cl_2 , $O_{0.3}$, M= 500.24, triclinic, P1, $\alpha = 8.6350(4)$ Å, b = 8.7930(3) Å, $c = 10.3560(6) \text{ Å}, \ \alpha = 82.257(3)^{\circ}, \ \beta = 73.341(2)^{\circ}, \ \gamma =$ 66.915(3)°, Z = 1, $D_{\text{calcd}} = 1.199 \text{ Mg} \cdot \text{m}^{-3}$, 2569 unique reflections, 2369 with $I > 2\sigma(I)$, $R_1 = 0.0445$ $[I > 2\sigma(I)]$ and $wR_2 = 0.1137$ [$I > 2\sigma(I)$], 302 parameters. All crystal data, fractional coordinates for the hydrogen and nonhydrogen atoms, equivalent thermal parameters and anisotropic temperature parameters for the non-hydrogen atoms, interatomic bond lengths and bond angles, and torsional angles have been deposited as Supporting Information.

IR Spectroscopy. IR spectra were scanned on an apparatus using a cell path length of 0.5 mm to investigate the amide N-H stretching (3200–3500 cm⁻¹), C=O stretching (amide I frequency at 1580–1720 cm⁻¹) and in-plane N-H bending (amide II frequency at 1480–1550 cm⁻¹). All the vinylog peptides also presented a weak to medium C=C stretching absorption at around 1640 cm⁻¹, but it turned out that it was not informative on the extended or folded conformational properties of the molecules. The peptide concentration was 0.005 M in DCM and in DMSO, and further dilution confirmed the absence of molecular aggregation. The N-H and C=O stretching frequencies were assigned on the basis of previous studies on related peptides and pseudopeptides and particularly on the influence of the adjacent groups.²¹

For this purpose, a Boc or Piv N-terminal group, and a Me or i-Pr C-terminal group attached to the same molecular backbone in some cases. For example, the NH frequency decreases by around 15 cm $^{-1}$ when changing NH-Me into NH-i-Pr as a C-terminal group. On the other hand, changing Piv into Boc as an N-terminal group decreases the CO stretching frequency by around 50 cm $^{-1}$ and increases the NH stretching frequency by around 20

cm $^{-1}$. In DCM, a free amide NH gives a sharp absorption in the $3400-3450~\rm cm^{-1}$ region and a free Piv-CO a strong absorption at $1610-1625~\rm cm^{-1}$. The occurrence of an N-H to C=O hydrogen bond results in a shift to low stretching frequencies for both NH and CO and to higher frequencies for the in-plane NH bending.

¹H NMR Spectroscopy. ¹H NMR spectra were run at 250 or 400 MHz with Me₄Si as internal reference, and the spin systems were solved by COSY and TOCSY experiments. The solvent accessibility, and therefore the extent of free or hydrogen-bonded character for the NH protons, was investigated in the pseudodipeptides by considering the shift of the NH proton resonance from $CDCl_3$ to $DMSO-d_6$, which proved to give more reliable results than the temperature coefficient for small model peptides. 21-23 The signal of a hydrogen-bonded, solventshielded NH is only slightly sensitive to the solvent, whereas the signal of a free, solvent-exposed NH is shifted downfield by DMSO- d_6 . In the case of a rapid equilibrium between an extended (solvent-exposed NH) and a folded (solvent-protected NH) conformer, the average solvent sensitivity of the NH resonance translates the relative percentages of both conformers.

Molecular Structures by X-ray Crystallography. The atomic structures of four *cis*-vinylog peptides and two trans-vinylog peptides have been solved by X-ray diffraction experiments. The crystal structures of Boc-"Procis"-NH-i-Pr 1d, Boc-"Protr"-NH-i-Pr 1c, Boc-"Protr"-Ala-NHi-Pr 5c, and Boc-"Alacis"-NH-i-Pr 1b have been published elsewhere. 10 The main torsional angles for the six vinylog derivatives are indicated in Table 2. The average dimensions of the vinylog fragment do not depend on the cis or trans disposition of the ethylenic bond (Figure 1). The geometry of the amide group is similar to that of the standard peptide group.²⁴ The ethylenic bond length is that of an isolated double bond, ²⁵ and both =C-C bonds have practically the same dimensions, suggesting that there is no electronic conjugation between the amide and the ethylenic bonds which are far from coplanarity (see angle ψ_2 in Table 2). As expected, the cis (Z) and trans (E) derivatives accommodate completely different conformations of the main chain. Both trans-derivatives 1c and 5c have an open structure with a cis-planar Boc-Pro amide bond and no intramolecular interaction. On the contrary, all four *cis*-vinylogs **1b**, **1d**, **2b**, and **8d** assume a folded conformation with an intramolecular NH to CO hydrogen bond closing a nine-membered pseudocycle that we propose to name the *cis*-vinylog turn.

All the amide groups are trans-planar. Figure 2 presents two examples of such open or folded crystal-lographic molecular structures, respectively **5c** and **8d**. A similar nine-membered folded conformation has been reported for the acylurea peptide *Z*-Val-Ni-Pr-CO-NH-*i*-Pr in the solid state, where the cis—trans urea fragment plays a role similar to the *cis*-ethylenic group.²⁶

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TABLE 2. Main Torsional Angles (deg) for the Six Crystallized Vinylog Peptides^a

$$\bigcup_{\substack{0 \\ \text{0} \\ \text{0}}}^{\text{N}} \bigvee_{\substack{\psi_1 \\ \text{0}}}^{\text{V}} \bigcup_{\substack{\psi_2 \\ \text{0}}}^{\text{0}} \bigcap_{\substack{\chi_2 \\ \text{0}}}^{\text{0}} \bigcup_{\substack{\chi_2 \\ \text{0}}}^{\text{0}} \bigvee_{\substack{\chi_2 \\ \text{0}}}^{\text{0}} \bigvee_{\substack{\chi_2 \\ \text{0}}}^{\text{0}} \bigcap_{\substack{\chi_3 \\ \text{0}}}^{\text{0}} \bigvee_{\substack{\chi_3 \\ 0}}^{\text{0}} \bigvee_{\substack{\chi_3 \\ 0}}$$

compound		ω_0	ϕ_1	ψ_1	ω_1	ϕ_2	μ_2	ν_2	ψ_2	ω_2	ϕ_3	ψ_3	ω_3
				cis-	Vinylog P	eptides							
Boc-"Pro ^{cis} "-NH- <i>i</i> -Pr	$\mathbf{1d}^b$				-174	-75	125	-3	-54	-177			
Boc-"Ala ^{cis} "-NH- <i>i</i> -Pr	$\mathbf{1b}^c$				172	-76	123	0	-57	-176			
Piv-"Ala ^{cis} "-NH- <i>i</i> -Pr	$\mathbf{2b}^d$				177	-82	122	2	-49	176			
Boc-Ala-"Pro ^{cis} "-Ala-NH-Me	$\mathbf{8d}^d$	173	-69	161	177	-72	125	-3	-53	-177	-65	139	173
standard <i>cis</i> -vinylog turn					180	-75	125	0	-55	180			
				trans	-Vinylog	Peptides	3						
Boc-"Pro ^{tr} "-NH- <i>i</i> -Pr	$\mathbf{1c}^b$				-6	-78	129	-177	-33	-179			
Boc-"Pro ^{tr} "-Ala-NH- <i>i</i> -Pr	$\mathbf{5c}^b$				1	-82	128	-179	-22	177	-85	141	177

^a The torsional angles are defined according to IUPAC-IUB Commission on Nomenclature. ^b See ref 10. ^c See ref 4e. ^d This work.

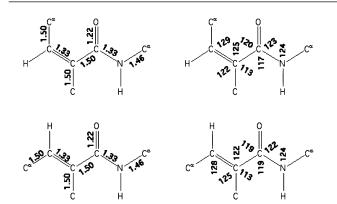


FIGURE 1. Average dimensions (Å, deg) for the *cis*- (up) and *trans*-vinylog (down) peptide unit in the crystal state.

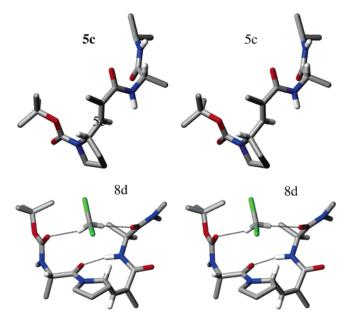


FIGURE 2. Stereoscopic view of the crystal molecular structures of the *trans*-vinylog dipeptide Boc-"Pro^{tr}"-Ala-NH-*i*-Pr **5c** and the *cis*-vinylog tripeptide Boc-Ala-"Pro^{cis}"-Ala-NH-Me **8d** with a dichloromethane molecule joining the N- and C-terminals.

Classically, the NH protons that are not involved in any intramolecular interaction are proton donors to a carbonyl of a neighboring molecule (Table 3). The *cis*-vinylog tripeptide Boc-Ala-"Procis"-Ala-NH-*i*-Pr **8d** mol-

ecule is also solvated by a water molecule, and the distance between the water oxygen and two amide oxygens is compatible with two O-H···O=C H-bonds. The dichloromethane molecule that crystallizes with 8d connects both N- and C-terminal carbonyls of the vinylog tripeptide skeleton, with two C to O distances that are a little shorter than a van der Waals contact (Figure 2).

Conformational Analysis in Solution. *cis*-Monovinylogs. In dichloromethane, all the *cis*-vinylog derivatives exhibit two components in the NH stretching absorption domain (Table 4). The high-frequency absorption at 3430–3455 cm⁻¹ classically corresponds to free NHs, while the low frequency domain at 3260–3290 cm⁻¹ is typical of H-bonded NHs. The invariant IR spectra with dilution suggesting the absence of molecular aggregation indicates that the *cis*-vinylog peptides are all folded by an intramolecular NH to CO H-bond that can be characterized by considering NH stretching frequency of Boc-"Procis"-NH-*i*-Pr 1d.

In **1d**, only the *i*-Pr-NH to Boc-CO H-bond is to be considered and forms what we name the *cis*-vinylog turn containing a nine-membered pseudocycle.

The low solvent sensitivity for derivatives **1b,d,f,h,j** and **2b,d** (-0.37 to 0.12 ppm) of the *i*-Pr-NH proton signal when changing from chloroform to DMSO supports the IR observation. In ¹H NMR spectroscopy, this observation still holds true for the other *cis*-vinylog derivatives where the acrylic NH resonance is systematically less solvent-sensitive than the other NH signals (Tables 5 and 6).

Another peculiarity of an H-bond is to shift the amide II IR absorption to high frequencies, and we effectively observe an absorption at around 1550 cm⁻¹ for the acrylamide, approximately 50 cm⁻¹ above the value for the N-terminal secondary amide (Table 4).

Insertion of an Ala residue at the C-terminus of the cis-vinylog ${\bf 1d}$ affords monovinylog dipeptides Boc-"Procis"-Ala-NH-R' ${\bf 5d}$ (R' = i-Pr) or ${\bf 6d}$ (R' = Me) which give rise to an additional NH stretching absorption at around 3340-3360 cm⁻¹, with medium intensity, which is attributed to the C-terminal NH on the basis of its variation with the adjacent Me or i-Pr group (Table 7). This observation is consistent with the rather low solvent effect (0.87 ppm) on the i-Pr-NH resonance of Boc-"Prot"-Ala-NH-i-Pr ${\bf 5c}$ in ${}^1{\rm H}$ NMR spectroscopy (Table 6).

The broad amide I profile prevents the assigning of the proton accepting carbonyl but, due to the very low occurrence of the γ -turn in dichloromethane for non Pro-



TABLE 3. Short X···O Distances (Å) in the X-H···O H-Bonds for the Six Crystallized Vinylog Peptides

		H-bond	X···O distance			
${\tt compound}^a$			intramolecular	intermolecular		
Boc-"Pro ^{cis} "-NH- <i>i</i> -Pr	$1d^a$	i-Pr-NH to Boc-CO	2.86			
Boc-"Alacis"-NH-i-Pr	$\mathbf{1b}^b$	$i ext{-Pr-NH to Boc-CO}$	2.86			
		"Alacis"-NH to "Alacis"-CO		2.92		
Piv-"Ala ^{cis} "-NH- <i>i</i> -Pr	$\mathbf{2b}^c$	<i>i</i> -Pr-NH to Piv-CO	2.86			
		"Alacis"-NH to "Alacis"-CO		3.00		
Boc-Ala¹-"Pro ^{cis} "-Ala²-NH-Me	$\mathbf{8d}^c$	Ala ² -NH to Ala ¹ -CO	2.84			
		Me-NH to Boc-CO		2.92		
		Ala¹-NH to Ala²-CO		2.94		
		WH to "Procis"-CO		3.18		
		WH' to Ala¹-CO		3.30		
		Cl_2CH_A to Boc-CO		3.31		
		$\mathrm{Cl_2CH_B}$ to $\mathrm{Ala^2\text{-}CO}$		3.22		
Boc-"Pro ^{tr} "-NH- <i>i</i> -Pr	$\mathbf{1c}^{a}$	$i ext{-Pr-NH to Boc-CO}$		3.02		
Boc-"Pro ^{tr} "-Ala-NH- <i>i</i> -Pr	$\mathbf{5c}^a$	$i ext{-Pr-NH to "Pro^{ ext{tr"}}-CO}$		2.84		
		Ala-NH to Boc-CO		3.04		

TABLE 4. IR Data for the Monovinylogs RCO-"Xaa"-NHi-Pr in Dichloromethane^a

		free I	NH	H-bonded NH	ar	nide I	ar	nide II
compound		"Xaa"-NH	i-PrNH	<i>i</i> -PrNH	Boc-"Xaa"	"Xaa"-CONH	Boc"Xaa"	"Xaa"-CONH
				cis-Derivative	S			
Boc-"Glycis"-NH-i-Pr	1h	3454		3284^{s}	1703	1667	1506	1548
Boc-"Ala ^{cis} "-NH- <i>i</i> -Pr	1b	3439		3283^{s}	1700	1666	1498	1549
Piv-"Ala ^{cis} "-NH- <i>i</i> -Pr	2b	3455		3252^{s}	1650	1663	1507	1552
Boc-"Valcis"-NH-i-Pr	1f	3440		3284^{s}	1697	1664	1496	1546
Boc-"Procis"-NH-i-Pr	1d		3443^{w}	$3260^{\rm s}$	1674	1665		1552
Piv-"Procis"-NH-i-Pr	2d		3443^{w}	$3242^{\rm s}$	1604	1664		1554
Boc-" β -Alacis"-NH- i -Pr	1j	3445	3434	3305^{w}	1706	1665		1508
				trans-Derivativ	es			
Boc-"Gly ^{tr} "-NH-i-Pr	1g	344	7		1711	1667		1505
Boc-"Ala ^{tr} "-NH- <i>i</i> -Pr	1a	344	.0		1709	1670		1502
Boc-"Val ^{tr} "-NH -i-Pr	1e	344	-1		1709	1669		1505
Boc-"Protr"-NH-i-Pr	1c		3445		1685	1674		1508
Piv-"Pro ^{tr} "-NH- <i>i</i> -Pr	2c		3450		1615	1668		1508
Boc-" β -Ala ^{tr} "-NH- i -Pr	1i	345	0	3377^{vw}	1709	1666		1508

TABLE 5. Chemical Shift δ and Solvent Sensitivity $\Delta\delta$ (ppm) in Chloroform/DMSO of the NH Proton Resonances for the cis-Monovinylog Peptides

$$\begin{array}{c|c}
O \\
N \\
-C^{\alpha} \\
-N \\
-C^{\alpha}
\end{array}$$

$$\begin{array}{c|c}
Me & H^3 & O \\
-N & C^{\alpha} \\
-N & C^{\alpha}
\end{array}$$

$$\begin{array}{c|c}
N \\
-M^4
\end{array}$$

	$ m NH^1$		NH^2		$ m NH^3$		$ m NH^4$	
	δ	$\Delta\delta$	δ	$\Delta\delta$	δ	$\Delta\delta$	δ	Δδ
1h			5.01/7.00	2.00	7.74/7.86	0.12		
1b			4.58/7.04	2.46	8.47/8.15	-0.32		
1f			4.59/7.05	2.46	8.29/8.11	-0.18		
1d					9.02/8.65	-0.37		
1j			5.03/6.84	1.81	5.59/7.59	2.00		
3b								
			7.14/8.08	0.94	8.59/8.50	-0.09		
			5.92/7.96	2.04	8.59/8.40	-0.19		
5d					9.45/9.05	-0.40	6.73/7.67	0.94
7d	5.34/6.96	1.62			9.23/9.34	0.11	6.57/7.66	1.09
	1b 1f 1d 1j 3b	δ 1h 1b 1f 1d 1j 3b	1h 1b 1f 1d 1j 3b	δ Δδ δ 1h 5.01/7.00 1b 4.58/7.04 1f 4.59/7.05 1d 1j 5.03/6.84 3b 7.14/8.08 5.92/7.96	$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$

containing related model peptides, 21 an $i+2\varnothing i$ H-bond involving the acrylic carbonyl is a highly improbable. Therefore, an $i+3\varnothing i$ H-bond with the Boc carbonyl and closing a 12-membered pseudocycle is the only possibility that is actually compatible with the cis-vinylog turn (Figure 3a).

Insertion of a Pro residue at the N-terminus of the *cis*-vinylog Boc-"Ala^{cis}"-NH-*i*-Pr **1b**, leading to the monovi-

nylog dipeptide Boc-Pro-"Ala^{cis}"-NH-*i*-Pr **3b**, induces less conformational perturbation. We only observe a very weak and broad absorption around 3330 cm⁻¹, which is attributed to the middle NH (Table 7) on the basis of the rather low solvent effect (0.94 ppm) on the NH proton resonance for the major trans form of the Boc-Pro amide bond (Table 5). This may denote a small percentage of γ -folded Pro residue, which is known to promote the



TABLE 6. Chemical Shift δ and Solvent Sensitivity $\Delta\delta$ (ppm) in Chloroform/DMSO of the NH Proton Resonances for the Trans Monovinylog Peptides

compound		$\mathrm{NH^{1}}$	$ m NH^1$		$ m NH^2$		$ m NH^3$		$ m NH^4$	
		δ	$\Delta \delta$	δ	$\Delta \delta$	δ	$\Delta \delta$	δ	$\Delta\delta$	
Boc-"Xaa ^{tr} "-NH- <i>i</i> -Pr										
"Gly ^{tr} "	1g			4.78/7.06	2.28	5.60/7.56	1.96			
"Ala ^{tr} "	1a			4.70/6.99	2.30	5.64/7.51	1.87			
"Val ^{tr} "	1e			4.54/6.88	2.34	5.52/7.50	1.98			
"Pro ^{tr} "	1c					5.54/7.55	2.00			
"β-Ala ^{tr} "	1i			4.67/6.86	2.19	5.63/7.43	1.80			
Boc-Pro-"Ala ^{tr} "-NH-i-Pr	3a									
trans-Boc-Pro				7.12/7.97	0.85	5.50/7.49	1.89			
cis-Boc-Pro				6.05/7.91	1.86	5.60/7.49	1.89			
Boc-"Pro ^{tr} "-Ala-NH- <i>i</i> -Pr	5c					6.25/7.62	1.37	6.75/7.62	0.87	
Boc-Ala-"Pro ^{tr} "-Ala-NH- <i>i</i> -Pr	7 c									
trans-Ala-"Pro ^{tr} "-		5.19/6.89	1.70			7.58/7.61	0.03	6.30/7.74	1.44	
cis-Ala-"Pro ^{tr} "-		5.38/6.99	1.61			6.41/7.52	1/11	5.91/7.66	1.75	

TABLE 7. IR Data for the cis- and trans-Monovinylog Dipeptides Boc-"Pro"-Ala-NH-R' and R-Pro-"Ala"-NH-R' in Dichloromethane^a

	$\rm free~NH_{\rm s}$		H-bond	H-bonded NHs		amide I			ide II	
compound				_						
			cis Deriv	atives						
Boc-"Procis"-AlaNH-i-Pr, 5d	345	$23^{b,c}$	$3230^{b,\mathrm{s}}$	$3340^{c, m}$		$1672^{d,e,f}$		1546^e	1524^f	
Boc-"Procis"-AlaNH-Me, 6d	3450^{c}		$3229^{b, w}$ $3362^{c, m}$			$1675^{d,e,f}$			$1550^{e,f}$	
Boc-Pro-"Ala cis "NH- i -Pr , ${f 3b}$	$3410^{b,\mathrm{m}}$	$3425^{c,\mathrm{w}}$	$3330^{b,\mathrm{w}}$	$3255^{c,\mathrm{s}}$	1695^d	1662^e	1672^{f}	1512^e	1551^{f}	
			trans Der	ivatives						
Boc-"Pro ^{tr} "-Ala-NH- <i>i</i> -Pr, 5c	349	$24^{b,c,\mathrm{s}}$	332	$25^{c,\mathrm{s}}$		$1682^{d,e,f}$		1500^e	$1524^{f,w}$	
Boc-"Protr"-Ala-NH-Me, 6c	$3429^{b,\mathrm{w}}$	$3449^{c,s}$	333	$37^{c,\mathrm{w}}$		$1684^{d,e,f}$		1500^e	$1543^{f,w}$	
Boc-Pro-"Alatr"-NH-i-Pr, 3a	3449^b	3418^c	330	$)5^{b,\mathrm{w}}$	1693^d	$1672^{e,f}$		15	$05^{e,f}$	
Piv-Pro-"Alatr"-NH-i-Pr, 4a	3445^b	3425^c	$3293^{b,\mathrm{w}}$		$3293^{b,\mathrm{w}}$ 1626^d $1670^{e,f}$		$1508^{e,f}$			

 a w, weak; m, medium; and s, strong absorption peak. b Ala-NH or "Ala"-NH. c R'NH. d RCO-Pro or RCO-"Pro". e Pro-CONH or "Pro"-CONH. f Ala-CONH or "Ala"-CONH.

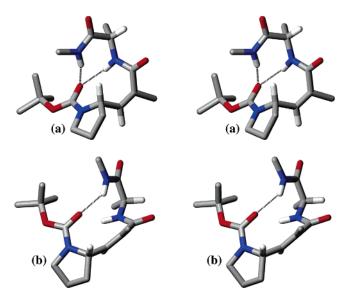


FIGURE 3. Stereoscopic view showing the $i+3\varnothing i$ H-bond between NH-Me and Boc carbonyl closing a 12-membered pseudocycle in (a) Boc-"Procis"-Ala-NH-Me **6d** and in (b) Boc-"Protr"-Ala-NH-Me **6c**.

 γ -turn in Pro-containing model peptides,²⁰ and this motif is compatible with an adjacent *cis*-vinylog turn. As a supporting argument, the middle NH resonance is very sensitive to solvent (2.04 ppm) for the minor cis form of

the Boc-Pro amide bond (Table 5), a conformation that is not compatible with the γ -turn.

When the cis-vinylog is in the middle position of a tripeptide, as for cis-Boc-Ala¹-"Procis"-Ala²-NH-i-Pr **7d** (Table 8), the NH stretching frequencies are quite similar to that for the monovinylog dipeptide Boc-"Procis"-Ala-NH-i-Pr **5d** (Table 7), with a more intense component at high frequencies due to free NHs. The NH protons also exhibit quite similar solvent accessibility (Table 5). We therefore conclude that the molecules share the same conformational properties. The great majority of the molecules **7d** accommodate the nine-membered cis-vinylog turn, and on the basis of the medium solvent sensitivity of the NH-i-Pr proton resonance (Table 5), a significant part of these molecules also contain an i-PrNH to Ala¹-CO H-bond closing a 12-membered pseudocycle (Figure 3a).

trans-Monovinylogs. The *trans*-monovinylogs RCO-"Xaa^{tr}"-NH-R' **1a**,c,e,g,i and **2a**,c only exhibit an NH absorption profile restricted to the high frequency domain in dichloromethane, while the Boc-"Xaa^{tr}" amide I absorption is around 10 cm⁻¹ higher than that for the cisanalogues, and the amide II absorption is observed at a lower frequency (Table 4). Moreover, the acrylamide NH proton resonance is more sensitive to solvation than for the *cis*-vinylog derivatives (Table 5). All these data



TABLE 8. IR Data for the cis- and trans-Monovinylog Tripeptides Boc-Ala¹-"Pro"-Ala²-NH-R' in Dichloromethane^a

	R'	free NHs	H-bond	led NHs	amide I			amide II			
7d 8d	<i>i-</i> Pr Me	3429 3444/3432	$3229^{b,\mathrm{s}} \ 3230^{b,\mathrm{s}}$	3339 ^{c,m} 3366 ^{c,m}	$\begin{array}{c} c \\ 1709^d \\ 1706^d \end{array}$	is - 1631^e 1633^e	1674^{f}	72 ^{f,g} 1683 ^g	$1502^{d,\mathrm{s}} \ 1501^{d,\mathrm{s}}$	1547 ^{f,m} 1549 ^{f,m}	$1525^{g, m w} \ 1530^{g, m w}$
7c 8c	<i>i</i> -Pr Me	3426 3434/3434	$3340^{b,\mathrm{s}}\ 3341^{b,\mathrm{s}}$		tro 1700^d 1697^d	$nns-1644^{e} \ 1640^{e}$	$1671 \\ 1671 \\ f$	$72^{f,g} \ 1684^g$	$1505^{d,\mathrm{s}} \ 1508^{d,\mathrm{s}}$	$1521^{f,\mathrm{m}} \\ 1525^{f,\mathrm{m}}$	$1505^{g, \rm s} \\ 1508^{g, \rm s}$

^a w, weak; m, medium; and s, strong absorption band. ^b Ala²-NH. ^c R'NH. ^d t-BuO-CONH. ^e CO-"Pro". f "Pro"-CONH. ^g CONHR'.

TABLE 9. IR Data for the Divinylog Dipeptides Boc-"Pro"-"Ala"-NH-i-Pr with Various Stereochemistry in Dichloromethane^a

compound		free NHs	H-bonded NHs	ami	ide I	amide II	
trans-trans derivative	9c 9g	$3447^{b,c,{ m s}} \ 3445^{b,c,{ m s,vw}}$	$3234^{b,c,\mathrm{vs}}$	1684^d 166	$1671^{e,f} \atop 8^{d,e,f}$		$\overline{5^{e,f,s}}$ $7^{e,f,s}$
trans-cis derivative cis-trans derivative	9d 9e	$3446^{b, m m} \ 3445^{c, m m}$	$rac{3256^{c,\mathrm{s}}}{3244^{b,\mathrm{s}}}$	$1685^d \\ 166$	$1668^{e,f} \\ 8^{d,e,f}$	$1504^{e} \ 1547^{e}$	$1549^f \\ 1508^f$

^a vw, very weak; m, medium;s, strong; and vs, very strong absorption band. ^b "Ala"-NH. ^c i-PrNH. ^d CO-"Pro". ^e "Pro"-CONH. ^f "-"Ala"-CONH.

indicate that the trans-vinylogs actually adopt extended conformations.

Addition of a Pro residue on the N-terminus in Boc-Pro-"Ala^{tr}"-NH-i-Pr 3a results in a weak and broad absorption band around 3305 cm $^{-1}$ attributed to Ala-NH in the present case (Table 7). The solvent sensitivity of the Ala-NH proton resonance depends on the cis or trans conformation of the Boc-Pro amide, and it is highly probable that trans-Pro is γ -folded by an $i+2\varnothing i$ H-bond while the trans-vinylog residue retains an extended conformation.

Similarly, the addition of an Ala residue at the C-terminus in Boc-"Prot"-Ala-NH–R′ $\mathbf{5c}$ and $\mathbf{6c}$ also results in a weak and broad absorption to appear around 3330 cm⁻¹, and its position depends on the Me or *i*-Pr C-terminal group. This is therefore assigned to a small percentage of H-bonded C-terminal NH which also shows a rather small solvent sensitivity (0.87 ppm, Table 6). Similarly, the broad amide I profile prevents the assigning of the carbonyl partner, but the very low occurrence of the γ -turn in dichloromethane for non Pro-containing related model peptides is to be noted, ^{4,21} and an H-bond involving the acrylic carbonyl in a γ -turn is not probable. Therefore, an H-bond with the Boc carbonyl closing a 12-membered pseudocycle is the only possibility (Figure 3b).

The two-component NH absorption profile for the trans-vinylog tripeptide Boc-Ala¹-"Protr"-Ala²-NH-i-Pr 7c is qualitatively similar to that for the trans-vinylog dipeptides Boc-"Protr"-Ala-NH-i-Pr 5c and Boc-Pro-"Alatr"-NH-i-Pr **3a**, but the low-frequency contribution is much more intense, indicating a higher percentage of intramolecular H-bonding. ¹H NMR reveals a cis-trans equilibrium of the Ala¹-Pro amide bond, and the major trans and the minor cis conformers have been characterized by the NOESY correlations between Ala¹-H^α and Pro-H^δ or Pro-H^α, respectively. In the major trans conformer, Ala² NH is highly protected from DMSO solvation with a low solvent variation of 0.03 ppm (Table 6) and is responsible for the absorption around 3340 cm⁻¹, which is actually not sensitive to the *i*-Pr \varnothing Me change (Table 8). The CO low stretching frequency of Boc-CO (which is

around 10 cm⁻¹ smaller than that for the *cis*-vinylog tripeptide **7c**) denotes an H-bonding participation, and the major trans conformer is therefore folded by an *i*-PrNH to Boc-CO interaction closing a 12-membered pseudocycle (Figure 3b). We may then conclude that the Ala-"Protr" sequence in *trans*-Boc-Ala-"Protr"-Ala-NH-*i*-Pr **7c** is more favorable to the above folded conformation than the Pro-"Ala^{tr}" sequence in *trans*-Boc-Pro-"Ala^{tr}"-NH-*i*-Pr **3a**. This observation is corroborated by the ¹H NMR structural analysis of various vinylogous hexapeptides containing the Val-"Protr" sequence which is actually folded by an NH to CO H-bond closing a 12-membered pseudocycle. ¹²

In the minor cis conformer of Boc-Ala¹-"Protr"-Ala²-NH-*i*-Pr, **7c**, Ala²-NH shows a rather low solvent sensitivity (Table 6) and is still engaged in an interaction with Boc-CO, also closing a 12-membered pseudocycle that is then flexible enough to accommodate a *cis*- or *trans*-ethylenic bond (Figure 3).

β-Alanine Vinylogs. Whatever the cis or trans disposition of the ethylenic bond, one additional carbon in the main chain of Boc-"β-Alacis"-NH-i-Pr 1j or Boc-β-"Alatr"-NH-i-Pr 1i results in a high solvent sensitivity for the i-Pr-NH resonance (Tables 5 and 6). However, the weak to very weak broad absorption at 3305/3377 cm⁻¹ in both cases (Table 4) indicates a small amount of folded molecules presenting a 10-membered pseudocycle. It is probably more flexible than the nine-membered pseudocycle in the cis-vinylog turn and is capable of accommodating a rigid trans-ethylenic bond to some extent although it affects only a small number of the molecules.

Divinylog Dipeptides. The spectroscopic observations on monovinylogs integrally hold true for molecules containing two consecutive vinylog residues. The high NH stretching frequencies, the low amide II frequencies (Table 9), and the high solvent accessibility (Table 10) for the *trans-trans*-divinylog dipeptide Boc-"Protr"-"Alatr"-NH-*i*-Pr **9c** coincide with the data for the *trans*-monovinylogs and denote an extended conformation devoid of any short-range intramolecular interaction. On the contrary, the low NH stretching frequencies, the high amide

TABLE 10. Chemical Shift δ and Solvent Sensitivity $\Delta\delta$ (ppm) in Chloroform/DMSO of the NH Proton Resonances for the Divinylog Dipeptides Boc-"Pro"-"Ala"-NH-i-Pr with Various Stereochemistry

		"Ala"-	NH	i-Pr-N	lΗ
compound		δ	$\Delta\delta$	δ	$\Delta\delta$
trans-trans-divinylog cis-cis-divinylog trans-cis-divinylog cis-trans-divinylog	9g 9d	5.71/7.83 9.40/9.07 5.70/7.80 9.38/8.92	-0.33 2.10	5.05/7.50 9.13/8.62 8.72/8.52 5.56/7.31	$\begin{array}{r} 2.45 \\ -0.51 \\ -0.20 \\ 1.75 \end{array}$

II frequencies (Table 9), and the low solvent accessibility (Table 10) for the cis-cis-divinylog dipeptide Boc-"Procis"-"Alacis"-NH-i-Pr **9g** coincide with the data for the cis-monovinylogs and denote two consecutive cis-vinylog turns (Figure 4). Both series of the above data are present for the cis-trans- and trans-cis-divinylog dipeptides Boc-"Procis"-"Alatr"-NH-i-Pr **9e** and Boc-"Protr"-"Alacis"-NH-i-Pr **9d**, so that each vinylog residue retains its own conformational properties in these molecules. Hence, we conclude that two consecutive vinylog residues are not capable of inducing a new structure containing a mediumrange H-bond differing from that already present in monovinylogs. We may therefore infer that an oligo cis-vinylog should adopt a helical structure with consecutive cis-vinylog turns.

Structural Influence of DMSO Solvation. All the above observations have been carried out in dichloromethane or chloroform, and we wonder whether the folded vinylog structures are stable in a strong solvating system such as DMSO. This solvent strongly solvates the amide NHs. It is also capable of breaking intramolecular H-bonding and therefore destabilizing folded structures.

The most informative IR region around the amide NH to amide CO intramolecular interactions in DMSO is the CO stretching absorption region (see Supporting Information), but because of the high dielectric constant, a small shift to lower frequencies by 2-5 cm⁻¹ with reference to dichloromethane must be taken into account. In ¹H NMR, the temperature coefficient for the NH proton resonances is commonly used for the structural analysis of peptides in DMSO. However, it has been reported that it can lead to erroneous conclusions for small model peptides.²² In this case, comparing the chemical shifts of analogous protons in related molecules seems to be of more valuable use. In the present case. the resonance of a free amide NH proton for the transmonovinylogs in DMSO is observed at 7.5-7.7 ppm (Table 6) and of a free urethane NH proton at 6.8-7.0 ppm. If an NH resonance is found at a higher chemical shift, it can be considered as protected from DMSO solvation by H-bonding.

The *trans*-monovinylogs Boc-"Xaa^{tr}"-NH-*i*-Pr in DMSO exhibit a single Boc-CO stretching frequency around 1710 cm⁻¹ ("Xaa" = "Gly" **1g**, "Ala" **1a**, "Val" **1e**) or 1685 cm⁻¹ ("Xaa" = "Pro" **1c**), a high frequency typical of an open conformer with a free Boc-CO. On the contrary, the Boc-CO stretching absorption for the *cis*-monovinylogs Boc-"Xaa^{cis}"-NH-*i*-Pr in DMSO exhibits two components denoting a conformational equilibrium. The high-frequency contribution (around 1705 cm⁻¹) corresponds to a free Boc-CO, and therefore to an open conformer, and the low-frequency contribution (around 1685 cm⁻¹ for "Xaa" = "Gly" **1h**, "Ala" **1b**, "Val" **1f**, and 1665 cm⁻¹ for

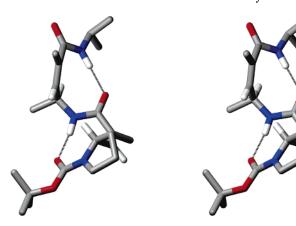


FIGURE 4. Stereoscopic view of the molecular structure associating two vinylog turns for the di-*cis*-vinylog dipeptide Boc-"Procis"-"Alacis"-NH-*i*-Pr **9g** in solution.

"Xaa" = "Pro" 1d) corresponds to an H-bonded Boc-CO, and therefore to a cis-vinylog folded conformer. The relative intensities of the two components are related to the populations of each conformer. Taking the intensity of the free component as a reference, "Procis" is the most favorable to folding in DMSO, with no evidence of open conformer, followed by "Valcis", "Alacis", and then "Glycis" with a *cis*-vinylog folding percentage of around 60, 50, and 30%, respectively. We also note that the i-PrNH proton resonance is lower in DMSO than in chloroform (Table 6). This observation still holds true for the cismonovinylog dipeptides Boc-"Procis"-Ala-NH-i-Pr 5d and Boc-Pro-"Alacis"-NH-i-Pr 3b (Table 5) and for the divinylog dipeptides containing one or two cis-residues (Table 10). We therefore conclude that the *cis*-vinylog turn has an intrinsic stability and is retained to a large extent in a strong NH-solvating medium.

We have pointed out that the folded structure presenting a 12-membered pseudocycle (Figure 3b) essentially concerns the trans-conformer of Boc-Ala-"Prot"-Ala-NH-*i*-Pr **7c**, and its retention in DMSO is questionable. In comparison with Boc-Ala-"Pro^{cis}"-Ala-NH-*i*-Pr **7d** where Boc-CO is free, the only support in favor of its partial retention in Boc-Ala-"Pro^t"-Ala-NH-*i*-Pr **7c** is the higher half width (28 versus 22 cm⁻¹) and the smaller CO stretching frequency (1703 versus 1706 cm⁻¹) of the Boc-CO amide I absorption. This suggests a two-component profile, the one at a higher frequency corresponding to the free Boc-CO, the other at a lower frequency denoting some amount of H-bonded Boc-CO. Hence, we conclude that the 12-membered pseudocycle is more sensitive to DMSO solvation than the nine-membered pseudocycle.

Conclusion

Insertion of an ethylenic group between the α-carbon and the carbonyl of a peptide residue results in a vinylogous peptide analogue also called vinylog. Starting from an α-amino aldehyde, it is rather easy to achieve either a *cis*- or *trans*-ethylenic bond and obtain the Boc-"Xaa^{cis}"-OH and Boc-"Xaa^{tr}"-OH synthons which can be classically introduced in a peptide chain or coupled with another vinylogous fragment. It is then possible to change one or several residues of a peptide into one or several vinylogous fragments. Moreover, the ethylenic bond may bear various substituting groups, allowing a chemical

modification on a second center beside the side chain in the starting α -amino aldehyde.

The structural consequences of inserting a vinylogous fragment in a peptide chain have been investigated for various sequences using X-ray diffraction in the solid state and ¹H NMR and IR spectroscopy in solution. The structural consequences greatly depend on the stereochemistry of this vinylogous fragment. A *cis*-vinylogous residue essentially promotes an NH to CO H-bond closing a nine-membered pseudocycle which stabilizes a folded motif that we propose to name the *cis*-vinylog turn. This *cis*-vinylog turn is particularly favored by the "Procis" residue, and it is stable enough to be retained in a strong solvating medium (DMSO). On the contrary, a *trans*-vinylogous residue essentially adopts an extended conformation.

A H-bond closing a larger pseudocycle containing 12 atoms is also observed in solution, but under our experimental conditions, it is only present in a small percentage of the molecules and particularly present in the Ala-"Protr" sequence but its retention in DMSO is questionable.

Two consecutive vinylogous residues retain their own structural preference, so that a "Xaa^{tr}"-"Xaa^{cis}" or "Xaa^{cis}"-"Xaa^{tr}" sequence is singly folded, and a "Xaa^{cis}"-"Xaa^{cis}" sequence is doubly folded. It seems to indicate that oligo vinylogs with all-cis, or all-trans, or alternating cis—trans stereochemistry could constitute new classes of foldamers presenting two centers for chemical modulation in each vinylogous residue. They could give access to additional tools for the rational design of potent peptide analogues.

Experimental Section

Vinylogous Aminoamides. General Procedure for the Preparation of the *trans*-Monovinylogous Aminoamides **1a, 1c, 1e, 1g, and 1i.** To a solution of *n*-BuLi (1.56 M in hexane, 9.22 mL, 14 mmol) in THF (20 mL) was added dropwise at -60 °C a solution of 2-diethylphosphonopropanoic acid²⁷ (1.43 g, 6.8 mmol) in THF (10 mL). After being stirred for 30 min at -60 °C, the aminoaldehyde 28 (6.8 mmol) in THF (5 mL) was added dropwise and the stirring was continued for 1 h at this temperature. The mixture was allowed to warm to room temperature during 30 min. After being stirred for an additional 3 h, it was hydrolyzed with water (20 mL). The organic layer was separated and washed with 10% aqueous $NaHCO_3$ (2 × 15 mL). The aqueous phases were combined and acidified to pH = 3.5 with 12 N HCl and extracted with Et₂O (2 × 30 mL). After drying over MgSO₄, the solvent was removed under vacuum to leave the crude carboxylic acid (60/ 40 < trans/cis < 98/2), which was purified by column chromatography. 14 To the so-obtained pure trans-carboxylic acid (0.70 mmol) dissolved in dichloromethane (10 mL) at room temperature were successively added triethylamine (0.071 g), (benzotriazol-1-yloxy)tris(dimethylamino)phosphonium hexafluorophosphate (BOP) (0.310 g, 0.70 mmol), followed after a few minutes by isopropylamine hydrochloride (0.070 g, 0.70 mmol) and triethylamine until basic pH was reached (around 0.100 g, 1.0 mmol). The resulting mixture was stirred at room temperature until TLC analysis indicated the total achievement (after around 2 h, depending to the substrate). The mixture was then diluted with dichloromethane (25 mL) and washed successively with 3 N HCl (3 \times 5 mL), brine (5 mL), aqueous saturated solution of NaHCO $_{\!3}$ (3 \times 5 mL), and brine (3 \times 5 mL). The organic layer was dried (MgSO $_{\!4}$) and concentrated. The resulting crude syrup monovinylogous aminoamide was purified by column chromatography on silica gel leading to pure trans isomer.

(4S)-(2E)-4-[[(1,1-Dimethylethoxy)carbonyl]amino]-2methyl-2-pentenoic Acid Isopropylamide (1a): 47.5 mg (88% isolated); white solid; mp 134 °C; R_f 0.32 (ethyl acetate/ hexane 1/1); IR (4.51 mM/DCM, CaF_2) ν_{max} 3440, 1709, 1630, 1670 cm $^{-1};$ $^{1}{\rm H}$ NMR (250 MHz, CDCl $_{3})$ $\delta_{\rm ppm}$ 1.15 (d, $^{3}J_{\rm H-H}$ = 6.5 Hz, 6 H), $1.19 \text{ (d, } {}^{3}J_{H-H} = 6.5 \text{ Hz}, 3 \text{ H}$), 1.41 (s, 9 H), 1.88 $(d, {}^{4}J_{H-H} = 1.0 \text{ Hz}, 3 \text{ H}), 4.08 \text{ (m, } {}^{3}J_{H-H} = 6.5 \text{ Hz}, 1 \text{ H}), 4.44$ (m, 1 H), 4.70 (d, ${}^{3}J_{H-H} = 7.0$ Hz, 1 H), 5.64 (d, ${}^{3}J_{H-H} = 7.0$ Hz, 1 H), 6.10 (d, ${}^{3}J_{H-H} = 8.5$ Hz, 1 H). ${}^{1}H$ NMR(250 MHz, C₆D₆) $\delta_{\rm ppm}$ 1.10 (d, ${}^3J_{\rm H-H}=6.6$ Hz, 3 H), 1.12 (d, ${}^3J_{\rm H-H}=6.5$ Hz, 6 H), 1.67 (s, 9 H), 1.98 (d, ${}^{4}J_{H-H} = 0.8$ Hz, 3 H), 4.40 (m, 1 H), 4.51 (d, ${}^{3}J_{H-H} = 6.8$ Hz, 1 H), 4.76 (m, 1 H), 5.33 (m, 1 H), 6.40 (d, ${}^{3}J_{H-H}$ = 8.1 Hz, 1 H). ${}^{1}H$ NMR (250 MHz, DMSO d_6) δ_{ppm} 1.09 (d, ${}^3J_{\text{H-H}} = 6.6 \text{ Hz}$, 6 H), 1.10 (d, ${}^3J_{\text{H-H}} = 6.3 \text{ Hz}$, 3 H), 1.38 (s, 9 H), $1.78 \text{ (d, } ^4J_{H-H} = 1.1 \text{ Hz, 3 H)}$, 3.94 (m, 1 H), $4.31~(\mathrm{m},~1~\mathrm{H}),~6.04~(\mathrm{dd},~^3J_{\mathrm{H-H}}=8.8~\mathrm{Hz},~^4J_{\mathrm{H-H}}=1.1~\mathrm{Hz},~1~\mathrm{H}),~6.99~(\mathrm{d},~^3J_{\mathrm{H-H}}=7.1~\mathrm{Hz},~1~\mathrm{H}),~7.51~(\mathrm{d},~^3J_{\mathrm{H-H}}=7.6~\mathrm{Hz},~1~\mathrm{H});$ ^{13}C NMR (62.896 MHz, CDCl₃) δ_{ppm} 12.9, 21.0, 22.7, 28.4, 41.5, $44.6, 79.5, 131.4, 136.8, 155.0, 168.0; [\alpha]_D -20.5^{\circ} (c 0.4, HCCl_3);$ MS (EI+) calcd for $C_{14}\ H_{26}N_2O_3\ [M]^+\ 270.2,$ found 271.2 [M +1]⁺. Anal. Calcd for C₁₄ H₂₆N₂O₃: C, 62.19; H, 9.69; N, 10.36. Found: C, 61.89; H, 9.52; N, 10.65.

General Procedure for the Preparation of the cis-Monovinylogous Aminoamides 1b, 1h, and 1j. n-BuLi (1.54 M in hexane, 1.54 mL, 2.37 mmol) was added dropwise to ethyl 2-diethylphosphonopropanoate (0,55 g, 2.37 mmol) in THF (10 mL) with being stirred at room temperature. After 20 min, the mixture was cooled to - 78 °C and the aminoaldehyde (Boc-Ala-H, Boc-Gly-H, or Boc- β -Ala-H, 2.26 mmol) in THF (10 mL) was added dropwise. After being stirred for 3 h, the reaction was quenched with an aqueous saturated ammonium chloride solution (12 mL) at -78 °C. The aqueous phase was extracted with Et₂O (3×30 mL), and the combined organic phases were washed with water $(2 \times 5 \text{ mL})$. Then, the organic layer was dried (MgSO₄), filtered, and concentrated under reduced pressure to afford a crude mixture of monovinylogous ester (60/40 \le cis/trans \le 90/10), which was purified by column chromatography on silica gel. To the so-obtained pure cis-vinylogous aminoester (1.20 mmol) in ethanol (6 mL) was added dropwise NaOH (1 N, 3 mL). The mixture was stirred at room temperature until TLC analysis indicated that all aminoester was reacted (2-3 h). Then HCl (1 N, 1.2 mL) was added, and the mixture was evaporated to remove ethanol. HCl (1 N, 1.6 mL) was added again, keeping the temperature below 25 °C, before the aqueous layer was extracted with EtOAc (3 \times 15 mL). The organic layers were combined and dried (MgSO₄). The solvent was removed under vacuum to afford the crude *cis*-monovinylogous amino acid. The coupling of this last compound with isopropylamine hydrochloride was similar to that described above to synthesize the transaminoamides 1a, 1c, 1e, 1g, and 1i. The crude product was purified on a silica gel chromatographic column leading to pure cis-aminoamide 1b, 1h, or 1j.

(4S)-(2Z)-4-[[(1,1-Dimethylethoxy)carbonyl]amino]-2-methyl-2-pentenoic Acid Isopropylamide (1b): 34.5 mg (84% isolated); crystals; mp 118 °C; R_f 0.43 (ethyl acetate/hexane 1/1); IR (4.86 mM/DCM, CaF₂) $\nu_{\rm max}$ 3439, 3283, 1700, 1666, 1636, 1549, 1498 cm⁻¹; ¹H NMR (250 MHz, CDCl₃) $\delta_{\rm ppm}$ 1.15 (d, ${}^3J_{\rm H-H}=6.5$ Hz, 6 H), 1.22 (d, ${}^3J_{\rm H-H}=6.5$ Hz, 3 H), 1.43 (s, 9 H), 1.90 (d, ${}^4J_{\rm H-H}=1.0$ Hz, 3 H), 4.12 (m, 1 H), 4.34 (m, 1 H), 4.58 (d, ${}^3J_{\rm H-H}=5.0$ Hz, 1 H), 5.13 (dd, ${}^3J_{\rm H-H}=10.5$ Hz, ${}^4J_{\rm H-H}=1.0$ Hz, 1 H), 8.47 (m, 1 H); ¹H NMR (250 MHz, C₆D₆) $\delta_{\rm ppm}$ 0.85 (d, ${}^3J_{\rm H-H}=6.7$ Hz, 3 H), 1.47 (d, ${}^3J_{\rm H-H}=6.5$

⁽²⁷⁾ Coutrot, P.; Snoussi, M.; Savignac, P. Synthesis 1978, 133–134.

⁽²⁸⁾ Boc-L-alanilal, Boc-L-β-alanilal, and Boc-L-valinal were prepared from corresponding Boc-L-amino acids according to Fehrentz and Castro. See: Fehrentz, J. A.; Castro, B. Synthesis 1983, 676–678. Boc-L-prolinal was synthesized similarly. See: Le Coz, S.; Mann, A.; Thareau, F.; Taddei, M. Heterocycles 1993, 36, 2073–2080. Some modifications were described by us to obtain the Boc-glycinal. See ref 14.

Hz, 3 H), 1.52 (d, ${}^3J_{\rm H-H} = 6.5$ Hz, 3 H), 1.58 (s, 9 H), 2.18 (d, ${}^4J_{\rm H-H} = 1.4$ Hz, 3 H), 4.32 (d, ${}^3J_{\rm H-H} = 6.0$ Hz, 1 H), 4.68 (m, 1 H), 4.72 (m, 1 H), 5.06 (dd, ${}^3J_{\rm H-H} = 10.3$ Hz, ${}^4J_{\rm H-H} = 1.4$ Hz, 1 H), 8.77 (m, 1 H); ${}^1{\rm H}$ NMR (250 MHz, DMSO- d_6) $\delta_{\rm ppm}$ 1.07 (d, ${}^3J_{\rm H-H} = 6.7$ Hz, 3 H), 1.10 (d, ${}^3J_{\rm H-H} = 6.6$ Hz, 3 H), 1.12 (d, ${}^3J_{\rm H-H} = 6.6$ Hz, 3 H), 1.39 (s, 9 H), 1.77 (d, ${}^4J_{\rm H-H} = 1.1$ Hz, 3 H), 3.91 (m, 1 H), 4.30 (m, 1 H), 5.06 (dd, ${}^3J_{\rm H-H} = 9.5$ Hz, ${}^4J_{\rm H-H} = 1.1$ Hz, 1 H), 7.04 (d, ${}^3J_{\rm H-H} = 6.7$ Hz, 1 H), 8.15 (d, ${}^3J_{\rm H-H} = 7.8$ Hz, 1 H). ${}^{13}{\rm C}$ NMR (62.896 MHz, CDCl₃) $\delta_{\rm ppm}$ 21.1, 21.2, 22.4, 28.3, 41.1, 45.6, 79.9, 131.1, 134.3, 155.8, 168.3; [α]_D +114.3° (c 0.6, HCCl₃); MS (FAB+) calcd for C₁₄ H₂₆N₂O₃ [M]+270.2, found 271.3 [M+1]+, 51%, 215.2, [M+1-t-Bu]+, 100%, 541.4, [2M+1]+, 55%. Anal. Calcd for C₁₄ H₂₆N₂O₃: C, 62.19; H, 9.69; N, 10.36. Found: C, 62.30; H, 9.45; N, 10.23.

General Procedure for the Preparation of the cis-Monovinylogous Aminoamides 1d and 1f. The ethyl bis-(trifluoroethyl)phosphonopropanoate²⁹ (1.08 g, 3.12 mmol) was added to a slurry of oil-free potassium hydride (0.125 g, 3.12 mmol) in THF (10 mL) at 0 °C. The solution was stirred for 15 min at this temperature followed by cooling to −78 °C. Boc-Pro-H or Boc-Val-H (2.6 mmol) in THF (5 mL) was added over 15 min. The reaction mixture was stirred at −78 °C for 3 h and then quenched with 2% aqueous HCl (15 mL). The aqueous phase was extracted with dichloromethane (3 × 50 mL). The organic layers were combined and dried (MgSO₄). After filtration, the solvent was removed under vacuum to afford the crude monovinylogous aminoester (cis/trans = 90/ 10) which was purified by column chromatography on silicagel. The so-obtained pure cis-monovinylogous aminoester was successively hydrolyzed into the corresponding cis-monovinylogous amino acid and coupled with isopropylamine hydrochloride as described above to give the crude corresponding monovinylogous aminoamide. The resulting syrup was purified on a silica gel chromatographic column leading to pure cismonovinylogous aminoamide 1d or 1f.

(2S)-2-[(1Z)-2-Methyl-3-[(1-methylethyl)amino]-3-oxo1-propenyl]-1-pyrrolidinecarboxylic Acid 1,1-Dimethylethyl Ester (1d): 10 34.0 mg (83% isolated); crystals; mp 150 °C; R_f 0.55 (ethyl acetate/hexane 1/1); IR (5.48 mM/DCM, CaF $_2$) $\nu_{\rm max}$ 3260, 1674, 1665, 1635, 1552 cm $^{-1}$, (6.36 mM/DMSO, CaF $_2$) $\nu_{\rm max}$ 1673, 1665, 1635, 1548 cm $^{-1}$; 1 H NMR(250 MHz, CDCl $_3$); 10 1 H NMR (250 MHz, DMSO- d_6) $\delta_{\rm ppm}$ 1.17 (d, $^{3}J_{\rm H-H}$ = 7.2 Hz, 6 H), 1.50 (s, 9 H), 1.51–2.27 (m, 4 H), 1.85 (s, 3 H), 3.31–3.48 (m, 2 H), 3.97 (m, 1 H), 4.57 (m, 1 H), 5.32 (d, $^{3}J_{\rm H-H}$ = 9.9 Hz, 1 H), 8.65 (d, $^{3}J_{\rm H-H}$ = 6.3 Hz, 1 H); 13 C NMR(62.896 MHz, CDCl $_3$) $\delta_{\rm ppm}$ 21.0, 22.4, 23.8, 28.4, 32.7, 41.0, 46.9, 51.4, 79.7, 129.0, 134.6, 155.0, 168.6; [α]_D +101.7° (c 0.6, HCCl $_3$); MS (EI+) calcd for C16 H28N2O3 [M]+ 296.4, found 297.1 [M+1]+. Anal. Calcd for C16 H28N2O3; C, 64.83; H, 9.52; N, 9.45. Found: C, 64.59; H, 9.69; N, 9.10.

General Procedure for the Preparation of the Monovinylogous Aminoamides 2a-d. A solution of pure cis- or trans-N-Boc-monovinylogous aminoamide 1a-j (3 mmol) in anhydrous Et₂O (20 mL) was submitted to a hydrochloride acid gaseous barbotage. A precipitate of monovinylogous aminoamide hydrochloride occurred before it dissolved in the reaction medium. When TLC analysis indicated the complete removal of Boc (around 2.5 h) and the complete formation of the corresponding monovinylogous aminoamide hydrochloride, the mixture was evaporated to give a paste. The so-obtained paste was washed with ether (5 \times 20 mL) before it was evaporated under vacuum to dryness. To this monovinylogous aminoamide hydrochloride (0.13 mmol) dissolved in CHCl₃ (4 mL) were added 2,2-dimethylpropanoyl chloride (0.024 g, 0.20 mmol) and diisopropylethylamine (DIPEA) (0.034 g, 0.26 mmol) at 0 °C. After being stirred for 1 h, the mixture was concentrated. The resulting crude cis- or trans-N-Piv-monovinylogous aminoamides 2a-d were purified on a silica gel chromatographic column.

(4S)-(2E)-4-(2,2-Dimethylpropionyl amino)-2-methylpent-2-enoic Acid Isopropylamide (2a): 64.0 mg (78% isolated); oil; R_f 0.28 (ethyl acetate/hexane 1/1); IR (4.25 mM/ DCM, CaF2) $\nu_{\rm max}$ 3460, 3250, 1670, 1550 cm $^{-1}$; $^1{\rm H}$ NMR (250 MHz, CDCl3) $\delta_{\rm ppm}$ 1.15–1.29 (m, 18 H), 1.95 (d, $^3J_{\rm H-H}=1.2$ Hz, 3 H), 4.00 (h, 1 H), 4.63 (m, 1 H), 5.48 (dd, 1 H, $^3J_{\rm H-H}=10.2$ Hz, $^4J_{\rm H-H}=1.2$ Hz, 1 H), 5.82 (m, 1 H), 8.85 (m, 1 H); $^{13}{\rm C}$ NMR (62.896 MHz, CDCl3) $\delta_{\rm ppm}$ 20.9, 22.4, 22.6, 27.5, 38.2, 41.0, 45.2, 130.0, 134.4, 168.1, 178.2; [α]_D -51° (c 0.3, HCCl3); MS (FAB+) calcd for C14 H26N2O2 [M]+ 254.2, found 255.3 [M+1]+, 100%. Anal. Calcd for C14 H26N2O2: C, 66.11; H, 10.30; N, 11.01. Found: C, 66.34; H, 10.65; N, 11.30.

Monovinylogous Dipeptides. General Procedure for the Preparation of the Monovinylogous Dipeptides 3a,b, 4a. The Boc protection of Boc-"Alati"-NH-i-Pr 1a or Boc-"Alacis"-NH-i-Pr 1b was removed by treatment with anhydrous HCl in Et₂O exactly as described above in the preparation of the monovinylogous aminoamides 2a-d. The procedure for the coupling of the so-obtained HCl, H-"Alati"-NH-i-Pr with Boc-Pro-OH or Piv-Pro-OH, leading to 3a or 4a, respectively, was the same as that employed above to prepare the vinylogous aminoamides 1a-j. This procedure was also used for the coupling between HCl, H-"Alacis"-NH-i-Pr, and Boc-Pro-OH that provided 3b.

 $(2S)\hbox{-}(E)\hbox{-}2\hbox{-}(3\hbox{-}Isopropylcar bamoyl-}1\hbox{-}methylbut-}2\hbox{-}enyl$ carbamoyl)-pyrrolidine-1-carboxylic Acid 1,1-Dimethyl **Ethyl Ester (3a):** 36.0 mg (87% isolated); white powder; mp 155 °C; R_f 0.49 (acetone/dichloromethane 3/7); IR (6.73 mM/ DCM, CaF $_2$) $\nu_{\rm max}$ 3449, 3418, 3305, 1693, 1672, 1505 cm $^{-1};$ (5.63 mM/DMSO, ${\rm CaF_2}$) $\nu_{\rm max}$ 1693, 1678, 1667, 1626, 1532 cm $^{-1};$ $^{1}{\rm H}$ NMR (250 MHz, CDCl₃) $\delta_{\rm ppm}$ 1.18 (d, $^{3}J_{\rm H-H}=6.5$ Hz, 6 H), 1.24 (d, ${}^{3}J_{H-H}$ = 6.8 Hz, 3 H), 1.47 (s, 9 H), 1.94 (d, ${}^{4}J_{H-H}$ = 0.7 Hz, 3 H, 1.81 - 2.50 (m, 4 H), 3.28 - 3.59 (m, 2 H), 4.07(m, 1 H), 4.17 (m, 1 H), 4.71 (m, 1 H), 5.60 (m, 1 H), 6.06 (d, ${}^{3}J_{H-H} = 7.5 \text{ Hz}, 1 \text{ H}), 6.05 \text{ (m, 1 H)}, 7.12 \text{ (m, 1 H)}; {}^{1}H \text{ NMR}$ (250 MHz, DMSO- d_6) $\delta_{\rm ppm}$ 1.09 (d, ${}^3J_{\rm H-H} = 6.6$ Hz, 6 H), 1.16 (d, ${}^{3}J_{H-H} = 6.5 \text{ Hz}$, 3 H), 1.34 (s, 9 H), (major), 1.40 (s, 9 H) (minor), 1.78 (m, 3 H), 1.65-2.28 (m, 4 H), 3.17-3.60 (m, 2 H), 3.81-4.19 (m, 2 H), 4.61 (m, 1 H), 6.13 (d, ${}^{3}J_{H-H} = 8.6$ Hz, 1 H) (minor), 6.15 (d, ${}^3J_{\rm H-H}=8.6$ Hz, 1 H) (major), 7.49 (d, ${}^3J_{\rm H-H}=7.4$ Hz, 1 H), 7.92 (d, ${}^3J_{\rm H-H}=7.7$ Hz, 1 H) (minor), 7.97 (d, ${}^3J_{\rm H-H}=7.7$ Hz, 1 H) (major); ${}^1{\rm H}$ NMR (250 MHz, DMSO- d_6) δ_{ppm} 1.09 (d, ${}^3J_{\text{H-H}} = 6.6$ Hz, 6 H), 1.16 (d, ${}^3J_{\text{H-H}} =$ 6.5 Hz, 3 H), 1.34 (s, 9 H) (major), 1.40 (s, 9 H) (minor), 1.78 $(\mathsf{m},\ 3\ \mathsf{H}),\ 1.65-2.28\ (\mathsf{m},\ 4\ \mathsf{H}),\ 3.17-3.60\ (\mathsf{m},\ 2\ \mathsf{H}),\ 3.81-4.19$ $(m, 2 H), 4.61 (m, 1 H), 6.13 (d, {}^{3}J_{H-H} = 8.6 Hz, 1 H) (minor),$ $6.15 \text{ (d, } ^3J_{H-H} = 8.6 \text{ Hz, } 1 \text{ H) (major)}, 7.49 \text{ (d, } ^3J_{H-H} = 7.4 \text{ Hz,}$ 1 H), 7.92 (d, ${}^{3}J_{H-H} = 7.7$ Hz, 1 H) (minor), 7.97 (d, ${}^{3}J_{H-H} =$ 7.7 Hz, 1 H) (major); 13 C NMR (62.896 MHz, CDCl₃) δ_{ppm} 13.0, 20.6, 22.6, 23.8, 24.4, 28.3, 27.5, 30.9, 41.5, 43.2, 47.0, 59.9, $61.1, 80.5, 132.4, 135.7, 155.8, 155.9, 168.1, 168.2, 171.3; [\alpha]_D$ -52° (c 0.3, HCCl3); MS (EI+) calcd for $C_{19}\,H_{33}N_3O_4$ [M]+ 367.5, found 368.2 $[M + 1]^+$. Anal. Calcd for C_{19} $H_{33}N_3O_4$: C, 62.10; H, 9.05; N, 11.43. Found: C, 62.15; H, 9.27; N, 11.25.

General Procedure for the Preparation of the Monovinylogous Dipeptides 5c,d and 6c,d. Monovinylogous dipeptide 5c or 6c was prepared by coupling Boc-"Pro"-OH (trans/cis = 80/20), obtained as described above in the synthesis of 1c and HCl, H-Ala-NH-i-Pr or HCl, H-Ala-NH-Me, respectively, using BOP as coupling reagent according to the procedure described above for the preparation of aminoamides 1a-j. Pure trans stereomer 5c or 6c was obtained after purification of the crude product (trans/cis = 80/20) on a silica gel chromatographic column. The same procedure was used to prepare 5d or 6d from Boc-"Pro"-OH (cis/trans = 90/10) obtained as described above in the synthesis of 1d and HCl, H-Ala-NH-i-Pr or HCl, H-Ala-NH-Me, respectively. Pure cis stereomer 5d or 6d was obtained after purification of the crude product (cis/trans = 90/10) on a silica gel chromatographic column.

(S)-(E)-(S)-2-{2-Methyl-3-[[1-methyl-2-[(1-methylethyl)-amino]-2-oxoethoxy] amino]-3-oxo-1-propenyl}-1-pyrro-

⁽²⁹⁾ Patois, C.; Savignac, P.; About-Jaudet, N.; Collignon, N. Synth. Commun. 1991, 21, 2391–2396.

lidine Carboxylic Acid 1,1-Dimethylethylester (5c): 14 45.0 mg (84% isolated); white solid; mp 114 °C; R_f 0.42 (ethyl acetate); IR (5.52 mM/DCM, CaF $_2$) $\nu_{\rm max}$ 3424, 3325, 1682, 1637, 1524, 1500 cm $^{-1}$; (5.64 mM/DMSO, CaF $_2$) $\nu_{\rm max}$ 1682, 1628, 1565 cm $^{-1}$; $^{1}{\rm H}$ NMR (250 MHz, CDCl $_3$), 13 $^{1}{\rm H}$ NMR(250 MHz, DMSO- d_6), $\delta_{\rm ppm}$ 1.03 (m, 6 H), 1.12 (d, $^{3}J_{\rm H-H}=7.0$ Hz, 3 H), 1.31 (s, 9 H), 1.80 (s, 3 H), 1.55–2.77 (m, 4 H), 3.17–3.45 (m, 2 H), 3.79 (m, 1 H), 4.28 (m, 1 H), 4.43 (m, 1 H), 6.15 (d, $^{3}J_{\rm H-H}=8.0$ Hz, 1 H), 7.62 (m, 2 H).

General Procedure for the Preparation of the Monovinylogous Tripeptides 7c-8d. Removal of Boc of monovinylogous dipeptides Boc-"Pro^{tr}"-Ala-NH-*i*-Pr 5c, Boc-"Pro^{cis}"-Ala-NH-*i*-Pr 5d, Boc-"Pro^{tr}"-Ala-NH-Me 6c, or Boc-"Pro^{cis}"-Ala-NH-Me 6d was effected with the procedure previously described using anhydrous gaseous HCl in Et₂O to afford HCl, H-"Pro^{tr}"-Ala-NH-*i*-Pr; HCl, H-"Pro^{cis}"-Ala-NH-*i*-Pr; HCl, H-"Pro^{tr}"-Ala-NH-Me; or HCl, H-"Pro^{tr}"-Ala-NH-Me; respectively. Then, these last compounds were coupled with Boc-Ala-OH using BOP as coupling reagent according to the procedure described above for the preparation of aminoamides 1a-j.

N-[(1,1-Dimethylethoxy)carbonyl]-L-alanyl-[(2E)-3-(pyrrolidin-2-yl)-2-methyl-2-propenoyl]-L-alanine Isopropy**lamide** (7c): 40.0 mg (74% isolated); powder; mp 194–195 °C; R_f 0.42 (acetone/ethyl acetate/hexane 1/1/1); IR (6.36 mM/DCM, CaF_2) ν_{max} 3426, 3340, 1700, 1672, 1644, 1521, 1505 cm⁻¹; (5.47 mM/DMSO, CaF $_2$) $\nu_{\rm max}$ 1704, 1664, 1655, 1636, 1533 cm $^{-1};$ ^{1}H NMR (250 MHz, CDCl $_{3})$ δ_{ppm} major trans conformer (Ala¹-Pro bond) 55% 1.11 (d, ${}^3J_{\rm H-H}=6.7$ Hz, 3 H), 1.15 (d, ${}^3J_{\rm H-H}=6.7$ Hz, 3 H), 1.28 (d, ${}^3J_{\rm H-H}=7.7$ Hz, 3 H), 1.43 (d, $^3J_{\rm H-H} =$ 7.0 Hz, 3 H), 1.43 (s, 9 H), 1.81–1.93 (m, 2 H), 1.95 $(s,\ 3\ H),\ 2.24-2.37\ (m,\ 2\ H),\ 3.48-3.76\ (m,\ 2\ H),\ 4.00\ (m,\ 1)$ H), 4.37-4.53 (m, 3 H), 5.20 (d, ${}^{3}J_{\rm H-H}=9.6$ Hz, 1 H), 6.11 (d, ${}^{3}J_{\rm H-H}=8.2$ Hz, 1 H), 6.30 (d, ${}^{3}J_{\rm H-H}=6.4$ Hz, 1 H), 7.58 (d, $^3J_{\mathrm{H-H}}=7.6$ Hz, 1 H); minor cis conformer (Ala¹-Pro bond) 45%: 1.14 (d, ${}^{3}J_{H-H} = 6.7 \text{ Hz}$, 3 H); 1.15 (d, ${}^{3}J_{H-H} = 6.7 \text{ Hz}$, 3 H), 1.30 (d, ${}^{3}J_{H-H} = 7.4$ Hz, 3 H), 1.38 (d, ${}^{3}J_{H-H} = 6.9$ Hz, 3 H), 1.43 (s, 9 H), 1.68–1.75 (m, 2 H), 1.95–2.18 (m, 2 H), 2.02 (d, $^4J_{\rm H-H}=0.9$ Hz, 3 H), 3.55–3.66 (m, 2 H), 4.02 (m, 1 H), 4.37-4.53 (m, 2 H), 4.80 (m, 1 H), 5.38 (d, ${}^{3}J_{H-H} = 7.8$ Hz, 1 H), 5.91 (d, ${}^{3}J_{H-H} = 7.6$ Hz, 1 H), 6.11 (d, ${}^{3}J_{H-H} = 8.2$ Hz, 1 H), 6.22 (d, ${}^{3}J_{H-H}$ = 9.7 Hz, 1 H); ${}^{1}H$ NMR (400 MHz, DMSO d_{6}) δ_{ppm} major trans conformer (Ala¹-Pro bond) 65% 1.04 (d, $^3J_{\mathrm{H-H}} = 7.0$ Hz, 3 H), 1.06 (d, $^3J_{\mathrm{H-H}} = 7.0$ Hz, 3 H), 1.17 (d, $^3J_{\mathrm{H-H}} = 6.5$ Hz, 3 H), 1.22 (d, $^3J_{\mathrm{H-H}} = 7.2$ Hz, 3 H), 1.38 (s, 9 H), 1.61 (m, 1 H), 1.86 (s, 3 H), 1.71-1.95 (m, 2 H), 2.03 (m, 1 H), 3.55–3.63 (m, 2 H), 3.81 (m, 1 H), 4.19–4.34 (m, 2 H), 4.66 (m, 1 H), 6.10 (d, ${}^3J_{\rm H-H}=8.7$ Hz, 1 H), 6.89 (d, ${}^3J_{\rm H-H}=7.2$ Hz, 1 H), 7.61 (d, ${}^3J_{\rm H-H}=7.3$ Hz, 1 H); 7.74 (d, ${}^3J_{\rm H-H}=7.6$ Hz, 1 H); minor cis conformer (Ala¹-Pro bond) 35%: 1.01 (d, $^3J_{\rm H-H}=7.0$ Hz, 3 H), 1.08 (d, $^3J_{\rm H-H}=7.0$ Hz, 3 H), 1.20 (d, $^3J_{\rm H-H}=7.0$ Hz, 3 H), 1.27 (d, $^3J_{\rm H-H}=7.0$ Hz, 3 H), 1.36 (s, 9 H), 1.61 (m, 1 H), 1.84 (s, 3 H), 1.71–1.95 (m, 2 H), 2.03 (m, 1 H), 3.55-3.63 (m, 2 H), 3.81 (m, 1 H), 4.13 (m, 1 H), 4.34 (m, 1 H), 4.66 (m, 1H), 6.19 (d, ${}^{3}J_{H-H}$ = 8.7 Hz, 1 H), 6.99 (d, ${}^{3}J_{H-H}$ = 7.2 Hz, 1 H), 7.52 (d, ${}^{3}J_{H-H}$ = 7.6 Hz, 1 H; 7.66 (d, ${}^{3}J_{H-H}$ = 7.4 Hz, 1 H); $^{13}\mathrm{C}$ RMN (62.896 MHz, CDCl₃) δ_{ppm} major trans conformer (Ala¹-Pro bond) 55%: 12.8, 16.6, 18.3, 22.6, 22.7, 24.9, 28.4, 33.4, 41.3, 46.9, 47.3, 49.4, 55.2, 80.0, 130.6, 135.4, 154.4, 168.0, 171.2, 171.7; minor cis conformer (Ala¹-Pro bond)-

45%: 13.4, 18.6, 18.9, 22.6, 24.9, 28.4, 30.9, 41.6, 46.8, 47.8, 49.0, 55.2, 79.5, 133.2, 136.3, 155.2, 169.0, 172.1, 172.4; $[\alpha]_D$ –90° (c 0.5, HCCl₃); MS (CI⁻) calcd for $C_{22}H_{38}N_4O_5$: 438.6 [M – 1]⁻, found 438.3, (49%), 364.4 [M – tBuO]⁻, 100%. Anal. Calcd for $C_{22}H_{38}N_4O_5$: C, 60.25; H, 8.73; N, 12.77. Found: C, 60.42; H, 8.48; N, 12.63.

General Procedure for the Preparation of the Divinylogous Dipeptides 9c-g and 10c-e. The divinylogous dipeptides 9c-g and 10c-e were obtained by coupling Bocor Piv-"Xaa^{tr}"-OH or Boc- or Piv-"Xaa^{cis}"-OH with the corresponding HCl, H-"Ala^{tr}"-NH-i-Pr or HCl, H-"Ala^{cis}"-NH-i-Pr. The coupling procedure was the same as that described for the preparation of aminoamides 1a-j.

 $(S)-(S)-2-\{2-Methyl-3-[[2(Z)-1,3-dimethyl-4-[(1-methyl$ ethyl)amino]-4-oxo-2-butenyl]amino]-(1E)-3-oxo-1-propenyl}-1-pyrrolidine Carboxylic Acid 1,1-Dimethylethyl **Ester (9e):** 52.0 mg (73% isolated); powder; mp 130 °C; R_f 0.32 (ethyl acetate); IR (4.29 mM/DCM, CaF_2) ν_{max} 3445, 3244, 1685, 1668, 1547, 1508 cm $^{-1}$; (4.29 mM/DMSO, CaF $_2$) $\nu_{\rm max}$ 1680, 1666, 1630, 1530 cm $^{-1}$; $^{1}{\rm H}$ NMR (250 MHz, CDCl $_3$) $\delta_{\rm ppm}$ 1.22 (d, ${}^{3}J_{H-H} = 6.5 \text{ Hz}$, 6 H), 1.26 (d, ${}^{3}J_{H-H} = 7.0 \text{ Hz}$, 3 H), 1.45 (s, 9 H), 1.90 (s, 3 H), 1.94 (s, 3 H), 1.52-2.13 (m, 4 H), $3.34 \text{ (m, }^{3}J_{H-H} = 7.0 \text{ Hz}, 2 \text{ H)}, 4.09 \text{ (m, 1 H)}, 4.51 \text{ (m, 1 H)},$ 4.76 (m, 1 H), 5.25 (d, ${}^{3}J_{\rm H-H}=10$ Hz, 1 H), 5.56 (d, ${}^{3}J_{\rm H-H}=7.5$ Hz, 1 H), 6.24 (dd, ${}^{3}J_{\rm H-H}=9.0$ Hz, ${}^{4}J_{\rm H-H}=1.0$ Hz, 1 H), 9.38 (d, ${}^{3}J_{H-H}$ = 6.5 Hz, 1 H); ${}^{1}H$ NMR (250 MHz, DMSO- d_{6}) $\delta_{\rm ppm}$ 1.13 (d, ${}^3J_{\rm H-H}$ = 6.5 Hz, 6 H), 1.20 (d, ${}^3J_{\rm H-H}$ = 6.5 Hz, 3 H), 1.43 (s, 9 H), 1.82 (s, 3H), 1.82 (s, 3H), 1.50–2.18 (m 4 H), 3.21-3.49 (m, 2 H), 3.91 (m, 1 H), 4.47 (m, 1 H), 4.64 (m, 1 H), 5.41 (d, ${}^{3}J_{H-H} = 10.75$ Hz, 1 H), 6.12 (d, ${}^{3}J_{H-H} = 9.0$ Hz, 1 H), 7.51 (d, ${}^{3}J_{H-H} = 7.0$ Hz, 1 H), 8.92 (d, ${}^{3}J_{H-H} = 6.5$ Hz, 1 H); 13 C NMR (62.896 MHz, CDCl₃) δ_{ppm} 13.0, 19.9, 21.0, 22.6, 23.7, 28.4, 32.6, 41.5, 43.8, 46.9, 55.3, 80.1, 130.3, 131.1, 133.5, 136.5, 155.2, 168.3, 168.8; $[\alpha]_D$ 57° (c 0.9, HCCl₃); MS (EI⁺) calcd for $C_{22} H_{37} N_3 O_4 [M+1]^+$: 407.27, found 407.0 (2%), 169.0 $[M - CH=C(CH_3)-CO-NH-CH(CH_3)-CH=C(CH_3)-CO-$ NH-iPr]⁺, 22%, 154.0 [M - $tBu-CO-NH-CH(CH_3)-CH=$ $C(CH_3)-CO-NH-iPr]^+$, 56%, 138.0 [M - $tBuO-CO-NH-CH(CH_3)-CH=C(CH_3)-CO-NH-iPr]^+$, 59%, 110 [M $tBuOCO-CO-NH-CH(CH_3)-CH=C(CH_3)-CO-NH-iPr]^+$ 100%. Anal. Calcd for $C_{22} H_{37} N_3 O_4$: C, 64.84; H, 9.15; N, 10.31. Found: C, 64.72; H, 9.40; N, 10.68.

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Supporting Information Available: General methods and description of the data for compounds 1c,e-j, 2b-d, 3b, 4a, 5d, 6c,d, 7d, 8c,d, 9c,d,f,g, and 10c,e. ¹H NMR (CDCl₃) of compounds 1a,b,e-j, 2b-d, 3a,b, 4a, 5c,d, 6c,d, 7d, 8c,d, 9c-g, and 10c-e. ¹H NMR (DMSO-d₆) of compounds 6c,d and 8d. ¹³C NMR of compounds 5c,d, 6c,d, 7d, 8c,d, 9c-g, and 10c-e. ¹H-¹H NMR (CDCl₃) of compound 7d. NOESY and TOCSY of compound 7c. Crystallographic information files (CIF format) for compounds 2b and 8d. This material is available free of charge via the Internet at http://pubs.acs.org.

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