α, ε -Hybrid Foldamers with 1,2,3-Triazole Rings: Order versus Disorder

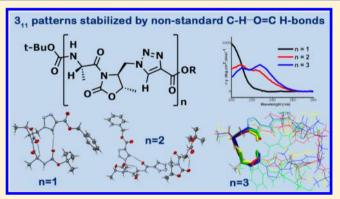
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Supporting Information

ABSTRACT: Two epimeric series of foldamers characterized by the presence of a repeating $\alpha_r e$ -dipeptide unit have been prepared and characterized by ¹H NMR and ECD spectroscopies together with X-ray diffraction. The first series contains L-Ala and D-4-carboxy-5-methyl-oxazolidin-2-one (D-Oxd). The other series contains L-Ala and L-Oxd. The L,D series of oligomers forms ordered β -turn foldamers, characterized by a 3_{11} pattern. The L,L series is not ordered. Simulations show that an ordered L,L trimer lies more than 2 kcal/mol higher than the more stable nonfolded extended conformations. Cu²⁺ forms complexes with both series but is not able to order the L,L series. Analysis of the EPR spectra shows that the L,D foldamers bear two types of complexation sites that are



assigned as a nitrogen donor of the triazole ring and a carboxylate ligand. The L-Ala-D-Oxd-Tri-CO motif may be introduced in any peptide sequence requiring the presence of a stable β -turn conformations, like in the study of protein-protein interactions.

INTRODUCTION

Pseudopeptide foldamers are synthetic molecules able to organize in well-defined secondary structures (i.e., helices, turns, and sheets). Despite their small size, they truly mimic biomacromolecules.¹ Their repetitive secondary structure is imparted by conformational restrictions of the monomeric unit.² Since 1996, when the neologism was coined, this research field has blossomed; many groups have explored oligomers with a wide backbone variety as potential foldamers, and several reviews have been published.³ In the past few years, we have extensively studied the conformational behavior of foldamers that contain the 4-carboxy-5-methyloxazolidin-2-one (Oxd) moiety.⁴ Acylation of the Oxd units yields imides that behave like rigid spacers, and the relative conformation of the two carbonyls generates a trans conformation with respect to the adjacent peptide bond.⁵ Among them, we have also investigated hybrid foldamers, where the Oxd moiety is alternated with an α - or a β -amino acid. In this case, the relative configuration of the Oxd and of the alternating amino acid are of paramount importance. For instance, the L-Ala-D-Oxd series tends to form β -bend ribbon spirals, while the L-Ala-L-Oxd series does not.⁶

Important stimulation came from the suggestion to employ dipeptide repeat units of various homologous amino acids.⁷ α / β -Hybrid peptides containing α - and β -amino acid constituents in a 1:1 backbone alternation displayed a variety of helical

structures.⁸ Some of these peptides elicited substantial biological activity.⁹ The concept was successfully extended to α/γ -, α/δ -, and β/γ -hybrid peptides.¹⁰ Recently, we have developed the synthesis of new preorganized dipeptide units that contain a 1,2,3-triazole (Tri) ring close to the Oxd group (Figure 1) and that mimic a constrained α,β -dipeptide

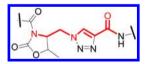


Figure 1. General structure of the Oxd-Tri scaffold. The preferential conformation of the imidic bond is shown. This scaffold is a constrained mimic of a ε -amino acid.

moiety.¹¹ The presence of the two heterocycles imparts a rigid *trans* conformation to the chain. The two rings differ for their spatial arrangement in that Oxd is chiral and 3,4-disubstituted while Tri is flat, achiral, and 1,4-disubstituted. The nitrogen of the Tri moiety cannot form a hydrogen bond, and the α , β -dipeptide moiety may be considered as a single unit, which constitutes a constrained mimic of ε -amino acid.¹²

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The Tri moiety may have interesting biological applications. For instance, pseudopeptides containing the Tri moiety have been successfully used to inhibit the activity of HIV-protease,¹³ as 1,2,3-triazoles efficiently mimic the amide bond of proteins. For the same reason, they have been successfully applied to the construction of artificial hybrid helices.¹⁴

In addition, the ability of triazole-based pseudopeptide structures to interact with metal ions (e.g., copper, zinc, etc.) provides the interesting possibility to exploit these systems as building blocks for supramolecular materials,¹⁵ as well as metal carriers¹⁶ or metal-stabilized folded structures.¹⁷

Figure 2 shows the structures of the two series of oligomers synthesized and studied in this work. The first series contains L-

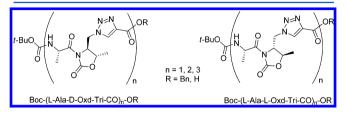


Figure 2. General formula of the oligomers investigated.

Ala and D-Oxd. The other series contains L-Ala and L-Oxd. The two epimeric series of foldamers characterized by the presence of the repeating α , ε -dipeptide unit can be considered prototypes for the investigation of the influence of asymmetric carbons and heterocycles on the secondary structure. The tendency to form stable complexes with Cu²⁺ ions was also assessed.

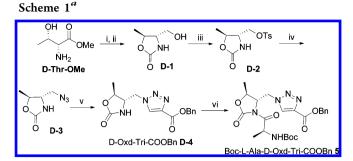
RESULTS AND DISCUSSION

1. Synthesis. The preparation of the L-Oxd-Tri-CO moiety has been recently described by our group.¹¹ It entails the formation of an intermediate azide derivative that is subsequently submitted to copper-catalyzed azide—alkyne cycloaddition (CuAAC) to afford the 1,2,3-triazole ring.¹⁸ In the course of this work, the synthesis of the key intermediates D- and L-4-(azidomethyl)-5-methyloxazolidin-2-one (D-3 and L-3) on a multigram scale was optimized in terms of yield, times, and costs. As compared with known procedures,¹⁹ a convenient one-pot process was developed for the preparation of (4S,5 S)-4-(hydroxymethyl)-5-methyloxazolidin-2-one D-1.

Briefly, D-**Thr-OMe** was cyclized with triphosgene at room temperature to give (4R,5S)-4-methylcarboxylate-5-methyloxazolidin-2-one (Scheme 1). After removal of the volatiles, the methyl ester was directly reduced to D-1 with sodium borohydride and converted in the tosylate D-2 with high yield. Finally, the conversion of D-2 to the corresponding azide D-3 was complete in 25 min under microwave irradiation with an overall yield of 79%.

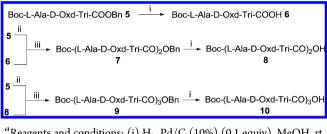
The azide D-3 was transformed into the corresponding 1,2,3triazole by CuAAC with benzyl propiolate, affording D-Oxd-Tri-COOBn D-4 in 78% yield.²⁰ The compound was coupled with Boc-L-Ala-OH to synthesize Boc-L-Ala-D-Oxd-Tri-COOBn 5. The latter was used to generate the corresponding dimer Boc-(L-Ala-D-Oxd-Tri-CO)₂-OBn 7 and trimer Boc-(L-Ala-D-Oxd-Tri-CO)₃-OBn 9 after deprotections and coupling reactions, as shown in Scheme 2.

Following the same approach, we prepared a series of oligomers that have the L_{JL} configuration to compare their conformational preferences with the L_{JD} series. Thus, L-Oxd-



^aReagents and conditions: (i) triphosgene (1.1 equiv), THF, rt, 1 h; (ii) NaBH₄ (1.1 equiv), EtOH, rt, 1 h, NH₄Cl, 45 min; (iii) TsCl (1.1 equiv), DMAP (10% w/w), pyridine, rt, 18 h; (iv) NaN₃ (1.1 equiv), dry DMF, MW 150 W/s, 25 min; (v) benzyl propiolate (1.0 equiv), DIEA (2.0 equiv), lutidine (2.0 equiv), CuI (0.1 equiv), dry acetonitrile, rt, 2 h; (vi) Boc-L-Ala-OH (1.0 equiv), HBTU (1.1 equiv), Et₃N (2.0 equiv), dry acetonitrile, rt, 50 min.

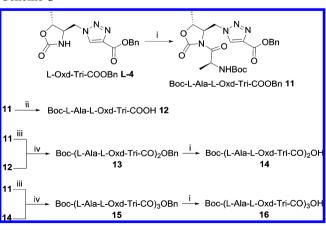
Scheme 2^a



"Reagents and conditions: (i) H_2 , Pd/C (10%) (0.1 equiv), MeOH, rt, 16 h; (ii) TFA (18 equiv), dry CH_2Cl_2 , rt, 4 h; (iii) HBTU (1.1 equiv), Et_3N (2.0 equiv), dry acetonitrile, rt, 50 min.

Tri-COOBn L-4 was prepared as previously described for D-4, starting from L-Thr-OMe, and it was further derivatized with Boc-L-Ala-OH to yield Boc-L-Ala-L-Oxd-Tri-COOBn 11 (Scheme 3). Similarly, the corresponding dimer Boc-(L-Ala-L-Oxd-Tri-CO)₂-OBn 13 and trimer Boc-(L-Ala-L-Oxd-Tri-CO)₃-OBn 15 were generated after deprotections and coupling reactions (Scheme 3).

Scheme 3^{*a*}



"Reagents and conditions: (i) Boc-L-Ala-OH (1.0 equiv), HBTU (1.1 equiv), Et₃N (2.0 equiv), dry acetonitrile, rt, 50 min; (ii) H_2 , Pd/C (10%) (0.1 equiv), MeOH, rt, 16 h; (ii) TFA (18 equiv), dry CH₂Cl₂, rt, 4 h; (iv) HBTU (1.1 equiv), Et₃N (2.0 equiv), dry acetonitrile, rt, 50 min.

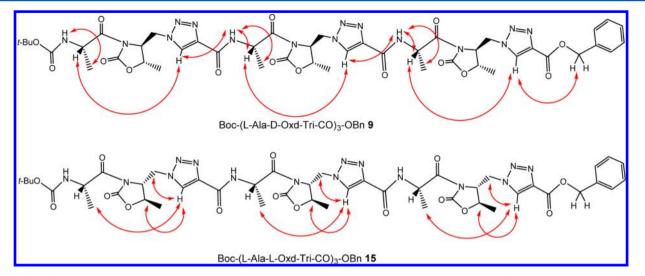


Figure 3. (Top) NOE enhancements gathered from the ROESY spectrum analysis of 9 (top) and of 15. (Bottom) (3 mM solution in CDCl₃, mixing time 0.400 s).

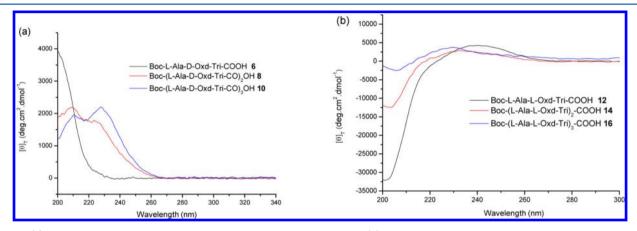


Figure 4. (a) Per-residue CD spectra of the oligomers of the D,L series: 6, 8, and 10. (b) Per-residue CD spectra of the oligomers of the L,L series: 12, 14, and 16.

2. Conformational Analysis. Preliminary information on the preferred conformation of the oligomers of both series was obtained by ¹H NMR and ECD absorption spectra in solution. These outcomes were further analyzed by single-crystal X-ray diffraction and molecular dynamics calculations. FT-IR spectroscopy in CH_2Cl_2 solution gave no useful information since we could not find any evidence for the formation of intramolecular N-H…O=C hydrogen bonds,²¹ as no stretching bands below 3400 cm⁻¹ were detected.

2.1. ¹H NMR spectroscopy. By ¹H NMR spectroscopy we confirmed the identity of all compounds. ROESY experiments performed on Boc-(L-Ala-D-Oxd-Tri-CO)₃-OBn **9** and Boc-(L-Ala-L-Oxd-Tri-CO)₃-OBn **15** gave information on the preferred conformation of the two oligomers. All experiments were recorded with a mixing time of 0.400 s for both samples (Figure 3).

Besides the trivial couplings, the ROESY spectrum of 9 in $CDCl_3$ shows cross peaks that have been attributed to the interaction of the triazole C–H with the amide NH and with the alanine CH (see the Supporting Information). The outcome suggests the formation of a preferred bent conformation. Unfortunately, **15** was poorly soluble in $CDCl_3$, which forced us to record the ROESY spectrum in CDc_3OD . The fast exchange between the NH of the amide

moieties and the deuterated solvent prevented the analysis of cross peaks that involve the amide NH, and we could record only the cross peaks between the triazole C-H and the alanine Me groups.

2.2. ECD Spectroscopy. To avoid the strong signal due to the benzyl groups, ECD spectroscopy was perfomed on the acids of both series, namely 6, 8, 10, 12, 14, and 16. In Figure 4 the per-residue ECD spectra of the oligomers of both series are reported. The spectra were recorded for 5 mM solutions in an 8:2 methanol/water mixture. The analysis of the preferred conformation of these compounds is not straightforward. These oligomers contain unusual structures that cannot easily be compared with α -amino acid oligomers. Nevertheless some information can be obtained from the comparison of these ECD spectra: the spectra of the L_JL series shown in Figure 4b are very similar and display a strong negative band at 200-204 nm and a broad positive featureless band between 220 and 260 nm. These ECD spectra are strongly reminiscent of random coils. By contrast, the ECD spectra of the L_D series (Figure 4a) differ from each other and suggest the appearance of a secondary structure in dimer 8 and trimer 10. Comparison with previous data,²² as well as the analogy between triazole and proline rings, suggests the assignment of the \sim 200 nm band to a β -turn that may form by hydrogen bonding between the

oxygen of the carbamate group (Boc) and the proton on the 1,2,3-triazole at position 5. The CD spectra of compounds 8 and 10 display two positive bands of comparable intensity at ~210 and ~228 nm, which recall an atypical helical structure. In fact, apart from the sign of the Cotton effect, these features are typical of right-handed helix- 3_{10} structures.²³ On the other hand, the ratio between the intensities of the two bands does not fully correspond to such structures. Taking into account the presence of a rigid triazole ring within the structure, by analogy with previous data, an atypical helix structure may be hypothesized, assuming that triazole is responsible for the inversion of the Cotton effect.

Based on the experimental evidence, we propose that the monomer, dimer, and trimer of the L,D series, i.e., compounds 6, 8, and 10, exhibit a β -turn responsible for the absorption at low wavelength. Chain elongation allows further structuring, stabilized by intramolecular hydrogen bonds.

2.3. Single-Crystal X-ray Diffraction. Crystals of 5 and 7. $0.5C_6H_6$ suitable for X-ray diffraction were grown as long needles by slowly evaporating solutions of 5 and 7 in benzene and benzene/methyl *tert*-butyl ether 1:1 v/v, respectively. Crystal data and refinement parameters for the two structures can be found in Table S1 (Supporting Information). In compound 5, the peptide folding mimics the β -turn structure of proteinogenic α -amino acids (Figure 5). The carbon atom C10

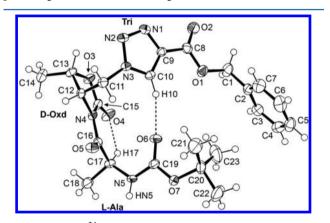


Figure 5. Ortep-3²⁴ plot of **5** with thermal ellipsoids drawn at 50% probability and H atoms represented as spheres of arbitrary radius. For simplicity, only H atoms engaged in intra- or intermolecular interactions are labeled. The dashed lines highlight intramolecular hydrogen bonding interactions discussed in the text.

of the triazole is engaged in a nonclassical hydrogen interaction with the carboxyl oxygen O6 of the Boc group $[d(H10\cdotsO6) =$ 2.14(2) Å], resulting in an 11-membered hydrogen-bonded turn (3₁₁-helical structure). By consequence, the two fivemembered rings in the D-Oxd-Tri residue are forced in a *syn* conformation around the C11–C12 linkage. At the same time, the *t*Bu carbon atom C23 of the Boc terminus is found at 3.51 Å from the center of the phenyl ring C2–C7, suggesting a CH/ π interaction. The α -carbon atom of the L-Ala residue (C17) has an intramolecular H-contact with the endocyclic carbonyl oxygen O4 of the D-Oxd fragment $[d(H17\cdotsO4) = 2.17(2) \text{ Å}].$

A similar pattern of intramolecular H-interactions, though with 0.2-Å longer H···A distances, is present in the C(O)-L-Ala2-D-Oxd2-Tri2 residue of $7.0.5C_6H_6$, which also exhibits a turn structure (Figure 6). Inspection of the backbone torsion angles for the two compounds (Table 1) indeed reveals a remarkable similarity between the conformation of L-Ala-D- Featured Article

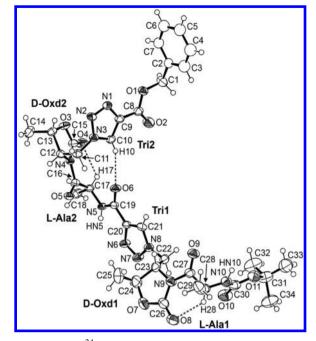


Figure 6. Ortep-3²⁴ plot of 7 with thermal ellipsoids drawn at 50% probability and H atoms represented as spheres of arbitrary radius. For simplicity, only H atoms engaged in intra- or intermolecular interactions are labeled. The dashed lines highlight intramolecular hydrogen bonding interactions discussed in the text.

Table 1. Selected Backbone Torsional Angles (in Degrees) for Compounds 5 and $7.0.5C_6H_6$

	5		7.0.5	7-0.5C ₆ H ₆	
C16-C17-N5-C19 (φ)	-60.1(3)	L-Ala	-57.2(3)	L-Ala2	
N4-C16-C17-N5 (<i>ψ</i>)	136.3(2)		147.2(2)		
C11-C12-N4-C16	71.5(2)	D-Oxd- Tri	72.2(3)	D-Oxd2- Tri2	
N3-C11-C12-N4	69.4(2)		64.8(3)		
N2-N3-C11-C12	90.1(2)		89.1(3)		
C27-C28-N10-C30 (ϕ)			-70.1(4)	L-Ala1-Tri1	
N9-C27-C28-N10 (<i>ψ</i>)			147.5(3)		
C22-C23-N9-C27			70.7(4)	D-Oxd1- Tri1	
N8-C22-C23-N9			172.6(3)		
N7-N8-C22-C23			-101.0(3)		

Oxd-Tri in monomer 5 and of L-Ala2-D-Oxd2-Tri2 in dimer 7; differences do not exceed 5° with the exception of the ψ angle for the L-Ala and L-Ala2 residues. The above-mentioned α -carbon/C=O contact is repeated also within the L-Ala1-D-Oxd1 unit (H28...O8), which however, does not adopt a turn structure. In fact, the two five-membered rings are rotated away from each other around the C22–C23 bond, as shown by the large N8–C22–C23–N9 torsion angle (172.6(3)). Moreover, D-Oxd1-Tri1 and D-Oxd2-Tri2 entail opposite rotations of the triazole moiety around the CH₂–N bond (N–N–C-C = $-101.0(3)^{\circ}$ vs 89.1(3)°). In the two compounds, the torsion angles are presented in Table 1. Overall, the torsion angles at the L-Ala residues (ϕ from -70.1 to -57.2° and ψ from 136.3 to 147.5°) are close to those typical for a PPII helix ($\phi = -79^{\circ}$ and $\psi = 150^{\circ}$).

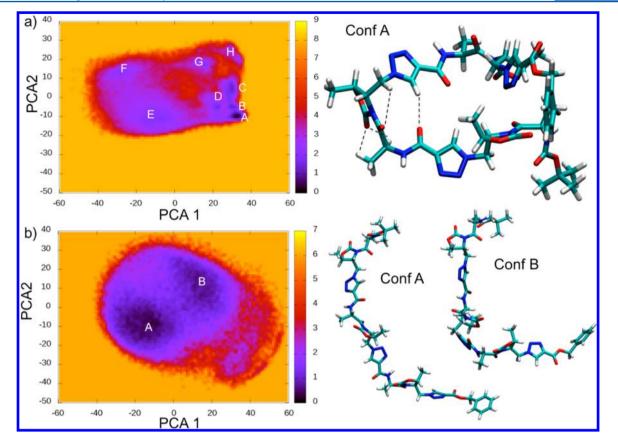


Figure 7. Conformational analysis of trimers 9 (a) and 15 (b). (Left) Two-dimensional free-energy landscape of the peptides (the values are in kcal mol^{-1}). The letters identify the most populated conformations. PCA1 and PCA2 are the two eigenvectors with the lowest eigenvalues of the principal component analysis calculated from the analysis of the MD trajectory. (Right) Representative structures of the conformational states corresponding to the most prominent minima of the free energy landscape.

Turning now to crystal packing, the carbonyl C16-O5 and the amide N5-HN5 groups in 5 are involved in intermolecular hydrogen bonds directed approximately parallel to the *b*-axis $[d(HN5\cdotsO5^{I}) = 2.15(2) A_{j}^{I} = x - 1/2, -y + 1/2, -z + 2].$ These two H-bonds per molecule link the peptides into supramolecular zigzag chains running along the a-axis. In the crystal lattice, molecules of 7 are linked by a H-bond between the amide (N5) and triazole (N1) nitrogen atoms, with $d(HN5...N1^{I}) = 2.138(17) \text{ Å } (I = x, y + 1, z).$ The second amide nitrogen N10 is also engaged in a H-bond with triazole nitrogen atom N7 of a neighboring molecule, but with a significantly longer distance $[d(HN10...N7^{II}) = 2.334(19) \text{ Å}; \text{ II}$ $= -x, y - \frac{1}{2}, -z$]. Both intermolecular contacts are directed approximately parallel to the b-axis of the unit cell. It is noteworthy that the only NH…CO hydrogen bonds found in these structures are of the intermolecular type and occur in 5, in agreement with the results of FT-IR spectroscopy in solution. Detailed geometrical parameters for hydrogen interactions are gathered in Tables S2 and S3 (Supporting Information).

2.4. Molecular Dynamics. The inability to grow crystals of the trimers 9, 10, 15, or 16 suggested the use of alternative approaches to gain insight in their structures. Molecular dynamics simulations for 1 μ s of the trimers Boc-(L-Ala-D-Oxd-Tri-CO)₃-OBn 9 and Boc-(L-Ala-L-Oxd-Tri-CO)₃-OBn 15 were carried out in an explicit methanol box, reproducing the conditions in which NMR spectra are registered. Principal component analysis (PCA) of the MD trajectories generated the free-energy landscapes of the two peptides (see the Experimental Section). This approach proved effective in reproducing the NMR and crystallographic data and provides insight into the free energy landscape.²⁵

The energy landscape of **9** is characterized by well-separated minima corresponding to specific conformational structures (Figure 7, see video V1 in the Supporting Information). The most populated conformer, Conf A, has a well-defined hydrogen-bond pattern that folds the peptide in a structurally stable turn structure. This 11-membered hydrogen-bonded turn corresponds to the motif identified in the solid state by the crystallographic analysis of the monomer **5** and dimer 7 (Table S4, Supporting Information). The other stable minima Conf B, Conf C, and Conf D maintain this structural motif, even if the other regions of the peptide show considerable conformational freedom (Figure 8).

This area of the peptide energy landscape, characterized by the presence of the turn, is around 2-7 kcal mol⁻¹ more stable than any other conformation (extended, helix-like, twist-like, Figure S1, Supporting Information). The nonstandard C-H··· O=C hydrogen bonds, represented by dashed lines in Figure 8, govern the structural stability of this motif.

On the contrary, in the energy landscape of **15** (Figure 8) the local minima are broad, and there is no significant basin of attraction indicative of a folded conformation. The most populated Conf A and Conf B (Figure 8) represent nonfolded extended conformations of the peptide that undergo practically free rotation in the solvent (video V2, Supporting Information). A turn structure can still be identified (Figure S2, Supporting Information). It is located the region of PCA1 that ranges

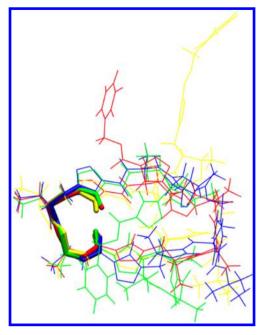


Figure 8. Superposition of the representative structures of the conformational states corresponding to Conf A (blue), Conf B (red), Conf C (yellow), and Conf D (green). In licorice, the conserved structural motif of the 11-membered hydrogen-bonded turn.

between 37 and 40, and its energy is more than 2 kcal/mol higher than the more stable minima. The β -turn structure of 9 is replaced by a γ -turn structure in 15 with a 7-membered hydrogen bonded ring. This structure usually competes with its 10-membered analogue (here 11 because of the presence of a ε amino acid) typical of a β -turn. Steric hindrance and/or chirality stabilize/destabilize the two competing ring-structures.

2.5. Complexation with Cu^{2+} . Triazole rings tend to interact with Cu^{2+} ions and form complexes. The extra stabilization energy provided by the interaction with Cu^{2+} could in principle be used to impart an ordered structure to both series of oligomers.

2.5.1. UV-vis Spectroscopy. UV-vis complexometric titrations show that oligomers 6, 8, and 10 interact with copper, giving rise to complexes. Table 2 presents the average

Table 2. $K_{\rm D}$ (mM) Values for the Cu²⁺ Complexes with 6, 8, and 10 in MeOH/H₂O 8:2 v/v^a

compd	λp (nm)	λt (nm)	$\binom{K_{\rm D}}{({ m mM})}$	Cu ²⁺ equiv at saturation
Boc-L-Ala-D-Oxd-Tri-COOH (6)	411	213	0.270	2
Boc-(L-Ala-D-Oxd-Tri-CO) ₂ - OH (8)	264	235	0.406	2-3
Boc-(L-Ala-D-Oxd-Tri-CO) ₃ - OH (10)	267	230	0.341	3

"The λ values corresponding to the peak and trough in the difference spectra as well as the Cu²⁺ equivalents at system saturation are reported.

 $K_{\rm D}$ values for the L_D series. The saturation threshold (i.e., the metal concentration corresponding to system saturation) increases with elongation of the pseudopeptide chain, due to the increase of putative metal-binding sites. The absence of absorption bands in the $\lambda \sim 600$ nm region suggests that we are dealing with type-II copper centers.

2.5.2. ECD Spectroscopy. Complexometric titrations of compounds 6, 8, and 10, monitored by ECD spectroscopy (Figure 9), showed that in all cases Cu^{2+} addition affects the intensity of the absorption bands without changing the spectral pattern. This suggests that copper leaves the folding unchanged as compared to the free ligand. This is supported by the lack of meaningful changes upon addition of an EDTA excess.

Analogous titrations with **12**, **14**, and **16** oligomers did not produce significant variations of the spectra (Figures S3–S5, Supporting Information). We conclude that Cu^{2+} is not able to induce extra folding in either series of oligomers.

2.5.3. EPR Spectroscopy. The interaction of Cu^{2+} with ligands 6, 8, and 10 in MeOH/H₂O 8:2 (v/v) was further characterized by EPR spectroscopy, and the corresponding spin-Hamiltonian parameters are summarized in Table 3. In all cases, two distinct complexation modes were highlighted: these correspond to at least two distinct metal binding sites, called a and b in Table 3.

The EPR parameters, with values of $A_{\parallel} > 100 \times 10^{-4} \text{ cm}^{-1}$, and the absence of absorption bands in the visible region of the electronic spectra suggest that, in all cases, the metal center is a type-II copper site. Center **a** shows lower A_{\parallel} values (106–109 $\times 10^{-4} \text{ cm}^{-1}$) as compared to center **b** (125–133 $\times 10^{-4} \text{ cm}^{-1}$). On the basis of these spectral parameters, center **a** was assigned to a nitrogen-coordinated copper (i.e., copper interacting with the triazole ring). Due to the similarity of its EPR parameters with those of $\text{Cu}(\text{H}_2\text{O})_6^{2+}$, center **b** was assigned to an oxygenated ligand (i.e., a carboxylate). $g_{\perp} < g_{\parallel}$ is typical of a distorted tetragonal geometry, with the unpaired electron located in the d_{x2-y2} orbital. The tetragonal distortion is further supported by the following evidence:

- $g_{\parallel}/A_{\parallel}$ ratios >222 × 10⁻⁴ cm⁻¹ for center **a** and >177 × 10⁻⁴ cm⁻¹ for center **b**. Square planar complexes exhibit much lower values ((105–135) × 10⁻⁴ cm⁻¹).²⁶ This also indicates that center **a** is more distorted than center **b**.
- The absence of electronic absorption bands in the visible region as, for complexes with strong tetragonal distortion, these bands shift toward the IR.²⁷

The lower degree of distortion of center \mathbf{b} as compared to center \mathbf{a} is consistent with its assignment to the carboxy-terminal site, since it is reasonable to expect a less conformationally constrained complex as compared to the triazole.

Finally, according to Sakaguchi et al.,²⁶ g_{\parallel} values >2.30 are consistent with predominantly ionic interactions and corroborate the assignments of binding sites **a** and **b** to a triazole and a carboxylate, respectively.

In all cases, the addition of a 10-fold molar excess of EDTA caused the formation of a pale blue solution. The EPR spectral pattern of the pale blue solutions was typical of Cu^{2+} -EDTA complexes. This indicates the complete disruption of the Cu^{2+} complexes by EDTA.

CONCLUSION

The purpose of this work was the optimized synthesis, conformational analysis, and application of two epimeric series of foldamers characterized by the presence of a repeating $\alpha_{,}\varepsilon_{-}$ dipeptide units. The ε -unit contains two heterocycles: a 1,2,3-triazole (Tri) and a 4-carboxy-5-methyloxazolidin-2-one (Oxd), while the α -unit is an alanine moiety. Both heterocycles imparts a rigid *trans* conformation to the chain, but the two rings differ

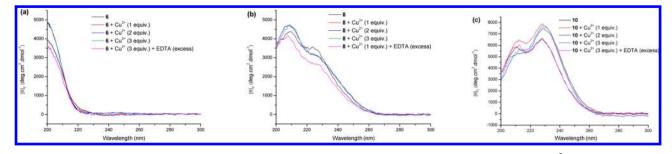


Figure 9. CD spectra of compounds 6, 8, and 10 dissolved in MeOH/H₂O 8:2 v/v, in the absence and presence of Cu^{2+} and after addition of a molar excess of EDTA (a) 6 (0.419 mM), (b) 8 (0.246 mM), and (c) 10 (0.174 mM).

Table 3. Spin-Hamiltonian Parameters Calculated from Simulation of the EPR Spectra of Cu^{2+} Complexes of Compounds 6, 8, and 10 Recorded in MeOH/H₂O 8:2 (v/v) at 77 K

ligand	center	contribution (%)	g_{\perp}	g∥	$A_{\parallel} \ (10^4 \ { m cm}^{-1})$	A_{\parallel} (G)	$g_{\parallel/A_{\parallel}}$ (cm)
6	a	82	2.085	2.427	106.5	113.9	227.88
	Ь	18	2.069	2.364	125.7	134.4	188.06
8	а	70	2.085	2.425	109.1	116.7	222.27
	ь	30	2.069	2.356	127.5	136.4	184.78
10	а	84	2.085	2.427	108.2	115.8	224.30
	Ь	16	2.070	2.362	133.3	142.6	177.19

for their spatial arrangement, as Oxd is chiral and 3,4disubstituted while Tri is flat, achiral, and 1,4-disubstituted.

Two series of oligomers have been prepared: Boc-(L-Ala-D-Oxd-Tri-CO)_n-OBn (L₂D series) and Boc-(L-Ala-L-Oxd-Tri-CO)_n-OBn (L₂L series). All compounds, including dimers and trimers, were prepared with good yields and have been characterized by ¹H NMR and ECD spectroscopies together with X-ray diffraction (when feasible).

Then both trimers were subjected to conformational analysis by atomistic molecular dynamics simulations. While the L,D series of oligomers form ordered β -turn foldamers, characterized by a 3₁₁ pattern, the L,L series is disordered. They also showed that an ordered L,L trimer lies more than 2 kcal/mol higher than the more stable non folded extended conformations.

Further information was obtained by tritration of both series with Cu^{2+} , which forms complexes with both series, but it is not able to order the L,L series.

Finally, analysis of the EPR spectra showed that the $L_{,D}$ foldamers bear two types of complexation sites that are assigned as the nitrogen of the triazole ring and a carboxylate ligand.

In conclusion, these results demonstrate that a small modification of the skeleton (D-Oxd versus L-Oxd) produces a dramatic variation in the properties of these new foldamers. While the L₁L series is basically unordered, even when complaxation with Cu²⁺ takes place, the L₁D-series forms in any case a β -turn conformation, characterized by a 3₁₁ pattern and stabilized by a nonconventional C-H…O=C hydrogen bond. This conformation is extremely stable, as it is present in all the oligomers of this series. Molecular dynamic calculations show that it is more stable of about 2–7 kcal/mol than any other.

These new foldamers containing the L-Ala-D-Oxd-Tri-CO motif may be introduced in any peptide sequence requiring the presence of a stable β -turn conformations, like in the study of protein—protein interactions.

EXPERIMENTAL SECTION

Synthesis. The melting points of the compounds were determined in open capillaries and are uncorrected. High-quality infrared spectra (64 scans) were obtained at 2 cm⁻¹ resolution using a 1 mm NaCl solution cell and a FT-infrared spectrometer. All spectra were obtained in 3 mM solutions in dry CH₂Cl₂ at 297 K. All compounds were dried in vacuo, and all of the sample preparations were performed in a nitrogen atmosphere. NMR spectra were recorded at 400 MHz (¹H NMR) and at 100 MHz (¹³C NMR). The measurements were carried out in CD₃OD and in CDCl₃. The proton signals were assigned by gCOSY spectra. Chemical shifts are reported in δ values relative to the solvent (CD₃OD or CDCl₃) peak.

(45,55)-4-(Hydroxymethyl)-5-methyloxazolidin-2-one (D-1). D-Thr-OMe (2.1 g, 12.6 mmol) was stirred in THF (15 mL) at room temperature until it was fully dissolved. Then triphosgene (3.9 g, 13.2 mmol) was added, and the reaction was stirred and refluxed for 1 h in an apparatus equipped with refrigerator and a trap containing CaCl₂ and NaOH. Then the mixture was cooled to room temperature, and the solvent was removed in vacuo. The (45,5S)-4-(carboxymethyl)-5methyloxazolidin-2-one was obtained in 86% yield (1.72 g) and directly transformed in D1.

Solid NaBH₄ (1.64 g, 11 mmol) was added to a stirred solution of (4*S*,5*S*)-4-(carboxymethyl)-5-methyloxazolidin-2-one (1.59 g, 10 mmol) in dry ethanol at 0 °C. After 5 min, the mixture was heated at room temperature and stirred for 1.5 h, and then a saturated aqueous solution of NH₄Cl (2.5 mL) was added and the mixture was stirred for additional 45 min. Then the mixture was filtered to eliminated the inorganic salts and concentrated in vacuo, leading to the formation of a waxy solid in 82% yield (1.08 g): $[\alpha]^{20}_{D} = -58.8 (c = 1.5, CHCl_3)$; IR (CH₂Cl₂, 3 Mm) ν 3611, 3452, 3280, 1760 cm⁻¹; ¹H NMR (CDCl₃, 400 MHz) δ 1.44 (3H, d, *J* = 7.2 Hz, Oxd-CH₃), 3.49–3.60 (2H, m, CHN-Oxd + CHNCHH), 3.65–3.79 (1H, m, CHN-CHH), 4.51 (1H, quintet, *J* = 5.6 Hz, CHO-Oxd), 6.73 (1H, bs, NH); ¹³C NMR (CDCl₃, 100 MHz) δ 20.3, 60.9, 62.8, 75.6, 160.3. Anal. Calcd for C₅H₉NO₃: C, 45.80; H, 6.92; N, 10.68. Found: C, 45.78; H, 6.91; N, 10.70.

[(45,55)-5-Methyl-2-oxooxazolidin-4-yl]methyl Methylparatoluensulfonate (p-2). To a stirred solution of p-1 (1.6 g, 12.6 mmol) and (dimethylamino)pyridine (0.13 g, 1.1 mmol) in pyridine (12 mL) was added tosyl chloride (3.4 g, 17.7 mmol) at 0 °C, and stirring was maintained for 18 h at room temperature.

After the end of the reaction, checked by TLC, CH_2Cl_2 (50 mL) was added, and the mixture was washed with brine (3 × 30 mL). The

organic phase was dried over sodium sulfate and concentrated under low pressure. The product was obtained pure after silica gel chromatography (cyclohexane/ethyl acetate 80:20 \rightarrow cyclohexane/ ethyl acetate 10:90 as eluant) in 74% overall yield as a white solid (2.66 g): mp = 95–98 °C; $[\alpha]^{20}_{\rm D} = -32.2$ (c = 1.5, CHCl₃); IR (CH₂Cl₂, 3 Mm): ν 3441, 1767, 1365 cm⁻¹; ¹H NMR (CDCl₃, 400 MHz) δ 1.38 (3H, d, J = 6.0 Hz, Oxd-CH₃), 2.43 (3H, s, Ar–CH₃), 3.64 (1H, dt, J = 5.6, 5.3 Hz CHN-Oxd), 3.96 (2H, dd, J = 5.3, Hz, CH₂-Ts), 4.37 (1H, quintet, J = 5.6 Hz, CHO-Oxd), 6,40 (1H, bs, NH), 7.24 (2H, d, J = 8.4 Hz, Ar), 7.75 (2H, d, J = 8.4 Hz, Ar); ¹³C NMR (CDCl₃, 100 MHz) δ 20.4, 21.7, 57.6, 69.4, 75.3, 127.8, 128.1, 130.0 130.3, 132.0, 145.5, 158.7. Anal. Calcd for C₁₂H₁₅NO₅S: C, 50.52; H, 5.30; N, 4.91. Found: C, 50.55; H, 5.28; N, 4.94.

(45,55)- 4-(Azidomethyl)-5-methyloxazolidin-2-one (*p*-3). To a stirred solution of compound *p*-2 (2.63 g, 9.2 mmol) in dry DMF (15 mL) was added sodium azide (0.66 g, 10.2 mmol). The mixture was stirred under microwave irradiation at 150 W for 25 min. At the end of the reaction, as checked by TLC, ethyl acetate (40 mL) was added to the mixture, and the organic phase was washed with water and with a 1 M aqueous solution of HCl. The organic layer was isolated, dried over sodium sulfate, and concentrated under low pressure. The product was obtained pure without any further purification in 79% yield (1.14 g) as a yellow oil: $[α]^{20}_{D} = -37.5$ (*c* 0.9, CHCl₃); IR (CH₂Cl₂, 3 Mm) ν 3440, 2120, 1764 cm⁻¹; ¹H NMR (CDCl₃, 400 MHz) δ 1.44 (3H, d, *J* = 6.4 Hz, Oxd-CH₃), 3.44 (2H, ABX, *J*_{AB} = 12.4 Hz, *J*_{AX} = 5.2 Hz, *J*_{BX} = 5.6 Hz, CH₂-N₃), 3.55 (1H, dd, *J* = 5.2, 12.0 Hz, CHN), 4.44 (1H, dq, *J* = 0.8, 6.8 Hz), 6.87 (1H, bs, NH); ¹³C NMR (CDCl₃, 100 MHz) δ 20.4, 53.5, 58.4, 76.3, 159.2. Anal. Calcd for C₅H₈N₄O₂: C, 38.46; H, 5.16; N, 35.88. Found: C, 38.42; H, 5.19; N, 35.85.

D-Oxd-Tri-COOBn (D-4). To a stirred solution of benzyl propiolate (0.25 g, 1.6 mmol) in dry acetonitrile (10 mL) were added diisopropylethylamine (3.2 mmol, 0.56 mL), lutidine (3.2 mmol, 0.37 mL), CuI (0.16 mmol, 0.03 g), and azide D-3 (1.6 mmol, 0.25 g) in this order at room temperature. Stirring was maintained for about 2 h at room temperature, and then, after the reaction was checked by TLC, the acetonitrile was removed under reduced pressure, replaced with ethyl acetate, and washed twice with brine $(3 \times 30 \text{ mL})$. The organic layer was isolated, dried over sodium sulfate, and concentrated under reduced pressure. The product was obtained pure after silica gel chromatography (cyclohexane/ethyl acetate $30:70 \rightarrow$ ethyl acetate 100% as eluant) in 78% yield (0.39 g) as a white solid: mp = 165-166°C; $[\alpha]_{D}^{20} = -52.0$ (c 1.2, CHCl₃); IR (CH₂Cl₂, 3 Mm): ν 3436, 3284, 1765 cm⁻¹; ¹H NMR (CDCl₃, 400 MHz) δ 1.36 (3H, d, J = 6.4 Hz, CHCH₃), 3.95 (1H, m, CHN), 4.41-4.55 (3H, m, CHO + CH_2CH), 5.38 (2H, AB, J = 12.8 Hz, CH_2Ph), 7.03 (1H, bs, NH), 7.25-7.47 (5H, m, Ph), 8.11 (1H, s, CH-triazole); ¹³C NMR (CDCl₃, 100 MHz) δ 20.2, 53.1, 58.5, 67.0, 75.7, 128.6, 128.7, 129.0, 135.2, 158.4, 160.3. Anal. Calcd for C₁₅H₁₆N₄O₄: C, 56.96; H, 5.10; N, 17.71. Found: C, 56.91; H, 5.14; N, 17.65.

Boc-L-Ala-D-Oxd-Tri-COOBn (5). A stirred solution of Boc-L-Ala-OH (0.09 g, 0.47 mmol) and HBTU (0.20 g, 0.52 mmol) in dry acetonitrile (20 mL) was stirred under nitrogen atmosphere for 10 min at room temperature. Then D-Oxd-Tri-OBn D-4 (0.150 g, 0.474 mmol) and DBU (0.141 mL, 0.948 mmol) in dry acetonitrile (10 mL) were added at room temperature. The solution was stirred for 2 h under nitrogen atmosphere, and then acetonitrile was removed under reduced pressure and replaced with ethyl acetate. The mixture was washed with brine $(1 \times 30 \text{ mL})$, 1 N aqueous HCl $(1 \times 30 \text{ mL})$, and with a concentrated solution of NaHCO₃ (1×30 mL), dried over sodium sulfate and concentrated in vacuo. The product was obtained pure after silica gel chromatography (cyclohexane/ethyl acetate 30:70 \rightarrow ethyl acetate 100% as eluant) as a white solid in 95% yield (0.22 g). Mp = 155–156 °C; $[\alpha]^{20}_{D}$ = –26.9; (*c* = 1.4, CHCl₃); IR (CHCl₃, 3 mM): ν 3440, 1789, 1704 cm⁻¹; ¹H NMR (CDCl₃, 400 MHz) δ 1.40 $(3H, d, J = 7.6 \text{ Hz}, \text{CHCH}_3), 1.42 (9H, s, t-Bu), 1.43 (3H, d, J = 6.0$ Hz, CHCH₃), 4.44 (1H, q, J = 3.6 Hz, CHN), 4.75–4.80 (2H, m, CH₂CH), 4.84 (1H, dq, J = 3.4, 5.6 Hz, CHO), 5.05 (1H, bs, NH), 5.27 (q, 1H, J = 6.8 Hz, CHAla), 5.41 (2H, AB, J = 13.6 Hz, CH₂Ph), 7.28-7.49 (5H, m, Ph), 8.78 (1H, s, CH-triazole); ¹³C NMR (CDCl₃, 100 MHz) δ 16.6, 20.5, 28.2, 49.2, 49.6, 59.7, 66.8, 73.5, 80.5, 128.3,

128.4, 128.5, 129.6, 135.5, 140.6, 151.3, 155.6, 160.2, 175.2. Anal. Calcd for $\rm C_{23}H_{29}N_5O_7:$ C, 56.67; H, 6.00; N, 14.37. Found: C, 56.70; H, 5.97; N, 14.35.

Boc-L-Ala-D-Oxd-Tri-COOH (6). Compound **5** (0.3 g, 0.61 mmol) was dissolved in MeOH (30 mL) under nitrogen. C/Pd (30 mg, 10% w/w) was added under nitrogen. A vacuum was created inside the flask using the vacuum line. The flask was then filled with hydrogen using a balloon (1 atm). The solution was stirred for 4 h under a hydrogen atmosphere. The product was obtained pure as an oil in 98% yield (0.24 g), after filtration through filter paper and concentration in vacuo: mp = 194 °C dec; $[\alpha]^{20}{}_{\rm D} = -21.9$ (c = 0.91, CHCl₃); IR 3440, 1789, 1709, 1705, 1700 cm⁻¹; ¹H NMR (CD₃OD, 400 MHz) δ 1.34 (3H, d, J = 6.0 Hz, CHCH₃), 1.36 (3H, d, J = 6.8 Hz, CHCH₃), 1.43 (9H, s, *t*-Bu), 4.51–4.58 (1H, m, CHN), 4.69–4.80 (3H, m, CH₂CH + CHO), 5.16–5.28 (m, 1H, CH-Ala), 8.60 (1H, s, CH-triazole); ¹³C NMR (CD₃OD, 100 MHz) δ 16.0, 19.2, 27.3, 49.2, 66.7, 74.2, 79.1, 128.8, 142.6, 152.1, 156.4, 174.7. Anal. Calcd for C₁₆H₂₃N₅O₇: C, 48.36; H, 5.83; N, 17.62. Found: C, 48.40; H, 5.85; N, 17.59.

Boc-(ι-Ala-D-Oxd-Tri-CO)₂OBn (7). A solution of 5 (0.14 g, 0.287 mmol) and TFA (0.14 mL, 1.80 mmol) in dry methylene chloride (20 mL) was stirred at room temperature for 4 h, then the volatiles were removed under reduced pressure and CF₃COO⁻ H₃N⁺-L-Ala-D-Oxd-Tri-COOBn was obtained pure without further purification as a waxy solid. ¹H NMR (CD₃OD, 400 MHz) δ 1.40 (3H, d, *J* = 6.4 Hz, CHCH₃), 1.53 (3H, d, *J* = 6.8 Hz, CHCH₃), 4.62–4.67 (1H, m, CH-Ala), 4.74–5.05 (4H, m, CH₂CH + CHO + CHN), 5.37 (2H, s, CH₂Ph), 7.29–7.40 (5H, m, Ph), 8.66 (1H, s, CH-triazole).

A solution of 6 (0.11 g, 0.287 mmol) and HATU (0.12 g, 0.32 mmol) in dry acetonitrile (20 mL) was stirred under nitrogen atmosphere for 10 min at room temperature. Then a mixture of the previously obtained CF3COO⁻ H3N⁺-L-Ala-D-Oxd-Tri-COOBn (0.143 g, 0.287 mmol) and DIEA (0.17 mL, 1.02 mmol) in dry acetonitrile (10 mL) was added dropwise at room temperature. The solution was stirred for 50 min under nitrogen atmosphere, then acetonitrile was removed under reduced pressure and replaced with ethyl acetate. The mixture was washed with brine (1 \times 30 mL), 1 N aqueous HCl (1 \times 30 mL), and with a concentrated solution of NaHCO₃ (1×30 mL), dried over sodium sulfate and concentrated in vacuo. The product was obtained pure after silica gel chromatography (c-Hex/ethyl acetate $30:70 \rightarrow$ ethyl acetate 100% as eluant) in 70% overall yield (0.16 g), as a white solid: mp = 164 °C; $[\alpha]^{20}_{D}$ = +48.5 (c = 1.0, CHCl₃); IR (CHCl₃, 3 mM) ν 3414, 1790, 1711, 1676 cm⁻¹; ¹H NMR (CDCl₃, 400 MHz) δ 1.37–1.48 (18H, m, CHCH₃ Ala +2 × CHCH₃ Oxd + t-Bu), 1.59 (3H, d, J = 7.2 Hz, CHCH₃ Ala), 4.40–4.50 (2H, m, 2 × CHN), 4.74-4.89 (6H, m, $2 \times CH_2$ CH + $2 \times CHO$), 5.21-5.31 (1H, m, CH-Ala), 5.41 (2H, s, CH₂Ph), 5.52-5.61 (1H, m, CH-Ala), 7.28-7.54 (6H, m, Ph + NH), 8.47 (1H, s, CH-triazole), 8.78 (1H, s, CHtriazole); $^{13}{\rm C}$ NMR (CDCl₃, 100 MHz) δ 16.5, 20.6, 28.2, 28.3, 29.7, 30.9, 37.5, 48.2, 48.7, 49.3, 59.2, 59.7, 73.4, 73.7, 74.0, 80.4, 127.5, 128.2, 128.4, 129.7, 135.5, 140.6, 142.7, 151.3, 159.7, 160.2, 173.6, 175.0. Anal. Calcd for $C_{34}H_{42}N_{10}O_{11}$: C, 53.26; H, 5.52; N, 18.27. Found: C, 53.29; H, 5.50; N, 18.31.

Boc-(*L*-Ala-*D*-Oxd-Tri-CO)₂-OH (**8**). For the synthetic procedure from 7 (0.77 g, 0.100 mmol), see the preparation of **6** given above (99% yield, 0.099 mmol, 0.67 g): mp = 200 °C; $[\alpha]^{20}_{D} = +20.5$ (*c* = 1.0, CH₂Cl₂); IR (CH₂Cl₂, 3 mM) ν 3446, 3412, 1790, 1718, 1670 cm⁻¹; ¹H NMR (CD₃OD, 400 MHz) δ 1.34–1.45 (18H, m, CHCH₃ Ala +2 × CHCH₃ Oxd + *t*-Bu), 1.50 (3H, d, *J* = 6.4 Hz, CHCH₃ Ala), 4.54–4.67 (2H, m, 2 × CHN), 4.74–4.95 (6H, m, 2 × CH₂CH + 2 × CHO), 5.20 (m, 1H, CH-Ala), 5.59 (m, 1H, CH-Ala), 8.47 (1H, s, CH-triazole), 8.78 (1H, s, CH-triazole); ¹³C NMR (CD₃OD, 100 MHz) δ 15.5, 15.6, 16.2, 27.3, 48.5, 49.2, 49.6, 51.2, 51.5, 52.4, 58.4, 59.6, 74.4, 75.9, 79.3, 127.8, 129.6, 142.2, 152.0, 152.1, 152.1, 156.4, 160.39, 160.5, 173.0, 173.2, 174.7. Anal. Calcd for C₂₇H₃₆N₁₀O₁₁: C, 47.93; H, 5.36; N, 20.70. Found: C, 47.88; H, 5.41; N, 20.74.

 $Boc-(L-Ala-D-Oxd-Tri-CO)_3-OBn$ (9). A solution of 5 (0.049 g, 0.100 mmol) and TFA (0.14 mL, 1.80 mmol) in dry methylene chloride (20 mL) was stirred at room temperature for 4 h, and then the volatiles were removed under reduced pressure and CF₃COO⁻ H₃N⁺-L-Ala-D-

Oxd-Tri-COOBn was obtained pure without further purification as a waxy solid.

A solution of 8 (0.068 g, 0.100 mmol) and HATU (0.042 g, 0.11 mmol) in dry acetonitrile (20 mL) was stirred under nitrogen atmosphere for 10 min at room temperature. Then a mixture of the previously obtained CF3COO⁻ H3N⁺-L-Ala-D-Oxd-Tri-COOBn and DIEA (0.04 mL, 0.3 mmol) in dry acetonitrile (10 mL) was added dropwise at room temperature. The solution was stirred for 50 min under nitrogen atmosphere, and then acetonitrile was removed under reduced pressure and replaced with ethyl acetate. The mixture was washed with brine $(1 \times 30 \text{ mL})$, 1 N aqueous HCl $(1 \times 30 \text{ mL})$, and a concentrated solution of NaHCO₃ (1 \times 30 mL), dried over sodium sulfate, and concentrated in vacuo. The product was obtained pure after silica gel chromatography (c-Hex/ethyl acetate $10:90 \rightarrow$ ethyl acetate 100% as eluant) in 51% yield (54 mg), as a white solid: mp = 194 °C; $[\alpha]^{20}_{D}$ = +69.0 (c = 0.97, CH₂Cl₂); IR (CH₂Cl₂, 3 mM) ν 3446, 3412, 1790, 1718, 1670 cm⁻¹; ¹H NMR (CDCl₃, 400 MHz) δ 1.38 (3H, d, J = 7.0 Hz, CH₃ Ala), 1.38 (3H, d, J = 7.0 Hz, CH₃ Oxd), 1.39 (9H, s, t-Bu), 1.43 (3H, d, J = 6.8 Hz, CH₃ Oxd), 1.46 (3H, d, J = 7.2 Hz, CH₃ Oxd), 4.41–4.52 (3H, m, 3 × CHN), 4.65–4.90 (9H, m, $3 \times CHO + 3 \times CH_2CH$), 5.17 (1H, d, J = 6.4 Hz, NH), 5.28 (1H, q, J = 6.4 Hz, CH Ala), 4.32-5.42 (2H, m, CH₂Ph), 5.52 (1H, m, CH Ala), 5.56 (1H, m, CH Ala), 7.29-7.39 (3H, m, Ph), 7.41-7.49 (2H, m, Ph), 7.57 (1H, d, J = 5.6 Hz, NH), 7.62 (1H, d, J = 5.6 Hz, NH), 8.50 (1H, s, CH-triazole), 8.64 (1H, s, CH-triazole), 8.75 (1H, s, CHtriazole); $^{13}{\rm C}$ NMR (CDCl₃, 100 MHz) δ 16.2, 16.6, 20.6, 28.2, 48.6, 48.8, 49.5, 59.6, 59.7, 59.8, 66.8, 73.9, 80.4, 127.8, 128.3, 128.5, 129.8, 135.5, 142.6, 151.3, 151.5, 155.9, 159.8, 173.6, 177.8. Anal. Calcd for C45H55N15O15: C, 51.67; H, 5.30; N, 20.09. Found: C, 51.72; H, 5.28; N, 20.11.

Boc-(*L*-Ala-*D*-Oxd-Tri-CO)₃-OH (10). For the synthetic procedure from 9 (0.05 mmol, 52 mg), see the preparation of 6 given above (99% yield, 0.049 mmol, 47 mg): mp = 206 °C; $[\alpha]^{20}_{D}$ = +120.0 (*c* = 0.90, CH₂Cl₂); IR (CH₂Cl₂, 3 mM) ν 3687, 3600, 3411, 1790, 1713, 1674, 1606, 1575, 1505 cm⁻¹; ¹H NMR (CDCl₃, 400 MHz) δ 1.22– 1.62 (27H, m, 3 × CH₃ Ala + 3 × CH₃ Oxd + *t*-Bu), 4.50–4.68 (3H, m, 3 × CHN), 4.69–4.99 (9H, m, 3 × CHO + 3 × CH₂CH), 5.14– 5.28 (1H, m, CH-Ala), 5.57–5.70 (2H, m, 2 × CH-Ala), 8.68 (1H, s, CH-triazole), 8.74 (1H, s, CH-triazole), 8.77 (1H, s, CH-triazole); ¹³C NMR (CDCl₃, 100 MHz) δ 15.7, 19.0, 27.3, 36.0, 37.6, 49.3, 59.6, 66.7, 74.1, 74.4, 79.2, 127.9, 142.2, 152.0, 156.5, 160.6, 173.4, 174.7. Anal. Calcd for C₃₈H₄₉N₁₅O₁₅: C, 47.75; H, 5.17; N, 21.98. Found: C, 47.70; H, 5.15; N, 22.02.

ι-*Oxd*-*Tri*-*COOBn* (*L*-4). For the synthetic procedure from L-3 (1.6 mmol, 0.25 g), see the preparation of D-4 given above (78% yield, 1.25 mmol, 0.39 g): mp = 167 °C; $[\alpha]^{20}_{D}$ = +53.0 (*c* 1.6, CHCl₃); IR (CH₂Cl₂, 3 Mm) ν 3436, 3284, 1765 cm⁻¹; ¹H NMR (CDCl₃, 400 MHz) δ 1.38 (3H, d, *J* = 6.4 Hz, CHCH₃), 3.95 (1H, m, CHN), 4.41–4.55 (3H, m, CHO + CH₂CH), 5.31 (2H, AB, *J* = 12.8 Hz, CH₂Ph), 7.03 (1H, bs, NH), 7.25–7.47 (5H, m, Ph), 8.26 (1H, s, CH-triazole); ¹³C NMR (CDCl₃, 100 MHz) δ 20.2, 53.1, 58.5, 67.0, 75.7, 128.6, 128.7, 129.0, 135.2, 158.4, 160.3. Anal. Calcd for C₁₅H₁₆N₄O₄: C, 56.96; H, 5.10; N, 17.71. Found: C, 56.91; H, 5.14; N, 17.65.

Boc-t-Ala-L-Oxd-Tri-COOBn (11). For the synthetic procedure from L-4 (1.2 mmol, 0.38 g), see the preparation of 5 given above (71% yield, 0.85 mmol, 0.41 g): mp = 117–118 °C; $[\alpha]^{20}_{D} = -38.0$ (*c* 0.8, CH₃OH); IR (CH₂Cl₂, 3 Mm) ν 3436, 1780, 1740, 1708 cm⁻¹; ¹H NMR (CDCl₃, 400 MHz) δ 1.38 (3H, d, *J* = 6.8 Hz, CHCH₃), 1.48 (12H, m, CH₃–Ala + *t*-Bu), 4.38 (1H, m, CHN), 4.65 (1H, d, *J* = 3.6, 6.4 Hz, CHO), 4.80 (2H, m, CH₂CH), 5.13 (1H, m, NH-Boc), 5.25 (1H, m, CH-Ala), 5.40 (2H, s, CH₂Ph), 7.25–7.43 (5H, m, Ph), 8.10 (1H, s, CH-triazole). ¹³C NMR (CDCl₃, 100 MHz) δ 15.2, 18.1, 20.2, 28.3, 49.6, 59.6, 67.1, 74.2, 128.5, 128.6, 135.1, 140.2, 151.3, 160.0, 175.0. Anal. Calcd for C₂₃H₂₉N₅O₇: C, 56.67; H, 6.00; N, 14.37. Found: C, 56.65; H, 5.99; N, 14.40.

Boc-*L*-Ala-L-Oxd-Tri-COOH (12). For the synthetic procedure from 11 (0.80 mmol, 0.32 g), see the preparation of 6 given above (99% yield, 0.79 mmol, 0.31 g): mp = 180 °C dec; $[\alpha]^{20}{}_{\rm D} = -48.7$ (*c* = 2.5, CHCl₃); IR (nujol): ν 3442, 3138, 1785, 1731, 1706, 1701, 1685 cm⁻¹; ¹H NMR (CD₃OD, 400 MHz) δ 1.32 (3H, d, *J* = 6.0 Hz, CHCH₃), 1.40 (9H, s, t-Bu), 1.43 (3H, d, J = 6.8 Hz, CHCH₃), 4.55– 4.59 (1H, m, CHN), 4.70–4.82 (2H, m, CH₂CH), 4.93 (m, 1H, CHO), 5.10–5.16 (m, 1H, CH-Ala), 8.32 (1H, s, CH-triazole); ¹³C NMR (CD₃OD, 100 MHz) δ 16.2, 18.9, 27.3, 29.3, 35.8, 49.7, 59.25 73.9, 74.2, 79.1, 129.1, 142.6, 152.1, 156.5, 174.7. Anal. Calcd for C₁₆H₂₃N₅O₇: C, 48.36; H, 5.83; N, 17.62. Found: C, 48.39; H, 5.80; N, 17.57.

*Boc-(ι-Ala-L-Oxd-Tri-CO)*₂-*OBn* (13). A solution of 11 (0.14 g, 0.287 mmol) and TFA (0.14 mL, 1.80 mmol) in dry methylene chloride (20 mL) was stirred at room temperature for 4 h, and then the volatiles were removed under reduced pressure and CF₃COO⁻ H₃N⁺-L-Ala-L-Oxd-Tri-COOBn was obtained pure without further purification as a waxy solid: ¹H NMR (CD₃OD, 400 MHz) δ 1.44 (3H, d, *J* = 6.4 Hz, CHCH₃), 1.52 (3H, d, *J* = 7.2 Hz, CHCH₃), 4.64 (1H, q, *J* = 3.8 Hz, CH-Ala), 4.73–4.81 (2H, m, CHHCH + CHN), 4.86–4.93 (1H, m, CHO), 4.98 (1H, dd, *J* = 4.0, 14.8 Hz, CHHCH), 5.35 (2H, AB, *J* = 13.2 Hz, CH₂Ph), 7.29–7.47 (5H, m, Ph), 8.64 (1H, s, CH-triazole).

A solution of 12 (0.113 g, 0.287 mmol) and HATU (0.12 g, 0.32 mmol) in dry acetonitrile (20 mL) was stirred under nitrogen atmosphere for 10 min at room temperature. Then a mixture of the previously obtained CF3COO⁻ H3N⁺-L-Ala-D-Oxd-Tri-COOBn and DIEA (0.17 mL, 1.02 mmol) in dry acetonitrile (10 mL) was added dropwise at room temperature. The solution was stirred for 50 min under nitrogen atmosphere, and then acetonitrile was removed under reduced pressure and replaced with ethyl acetate. The mixture was washed with brine $(1 \times 30 \text{ mL})$, 1 N aqueous HCl $(1 \times 30 \text{ mL})$, and a concentrated solution of NaHCO₃ (1 \times 30 mL), dried over sodium sulfate, and concentrated in vacuo. The product was obtained pure after silica gel chromatography (c-Hex/ethyl acetate $30:70 \rightarrow$ ethyl acetate 100% as eluant) in 70% overall yield (0.20 mmol, 0.15 g), as a white solid: mp = 110 °C; $[\alpha]^{20}_{D} = -60.5$; IR (CHCl₃, 3 Mm) ν 3438, 3434, 3410, 1674, 1791, 1743, 1710 cm⁻¹; ¹H NMR (CD₂OD, 400 MHz) δ 1.31 (3H, d, J = 7.2 Hz), 1.41 (9H, s, t-Bu), 1.43 (15H, m, CH₃-Oxd), 1.46 (3H, d, J = 8 Hz), 4.51-4.65 (3H, m, CHN-Oxd), 4.69-4.80 (4H, m, CHO-Oxd), 5.11 (1H, q, J = 6.8 Hz), 5.36 (2H, s), 5.55 (1H, q, J = 7.2 Hz), 7.27–7.47 (5H, m), 8.37 (1H, s, CHtriazole), 8.58 (1H, s, CH-triazole); ¹³C (CD₃OD, 100 MHz) δ 17.6, 20.3, 28.7, 29.5, 38.9, 48.3, 49.0, 49.6, 50.3, 51.0, 53.9, 59.7, 60.6, 67.9, 75.8, 77.3, 80.6, 129.0, 129.6, 131.0, 137.0, 140.7, 143.4, 153.5, 157.9, 161.6, 174.5, 175.8. Anal. Calcd for $C_{34}H_{42}N_{10}O_{11}\!\!:$ C, 53.26; H, 5.52; N, 18.27. Found: C, 53.22; H, 5.48; N, 18.26.

Boc-(*ι*-Ala-*L*-Oxd-Tri-CO)₂-OH (14). For the synthetic procedure from 13 (0.20 mmol, 0.15 g), see the preparation of 6 given above (99% yield, 0.19 mmol, 0.13 g): mp = 200 °C; $[\alpha]^{20}{}_{D} = -60.9$ (c = 1.0, CH₃OH); IR (nujol) ν 3373, 3144, 1781, 1744, 1707, 1663, 1577, 1507 cm⁻¹; ¹H NMR (CD₃OD, 400 MHz) δ 1.26–1.51 (21H, m, 2 × CHCH₃-Ala +2 × CHCH₃-Oxd + *t*-Bu), 4.52–4.82 (8H, m, 2 × CHN + 2 × CHO + 2 × CH₂CH) 5.11 (1H, q, *J* = 6.8 Hz, CH-Ala), 5.57 (1H, q, *J* = 6.8 Hz, CH-Ala), 8.28 (1H, s, CH-triazole), 8.40 (1H, s, CH-triazole); ¹³C NMR (CD₃OD, 100 MHz) δ 16.2, 16.3, 19–0 27.3, 48.9, 49.6, 52.3, 52.4, 58.3, 58.4, 58.5, 59.6, 74.4, 75.9, 79.2, 127.1, 127.6, 129.2, 142.0, 142.2, 152.1, 156.5, 159.3, 160.4, 173.0, 173.0, 173.2, 174.4. Anal. Calcd for C₂₇H₃₆N₁₀O₁₁: C, 47.93; H, 5.36; N, 20.70. Found: C, 47.94; H, 5.38; N, 20.73.

*Boc-(ι-Ala-L-Oxd-Tri-CO)*₃-*OBn* (**15**). For the synthetic procedure from **11** (0.15 mmol, 0.10 g), see the preparation of **9** given above (40% yield, 0.04 mmol, 42 mg): mp = 108 °C; $[\alpha]^{20}_{D} = -17.5$; IR (CHCl₃, 3 Mm) ν 3447, 3433, 3410, 1789, 1772, 1749, 1712, 1676 cm⁻¹; ¹H NMR: (CD₃OD): δ 1.31 (3H, d, *J* = 6,8 Hz, CH₃-Ala), 1.41 (9H, s, *t*·Bu), 1.48 (3H, d, *J* = 6 Hz, CH₃-Ala), 1.49 (3H, d, *J* = 7.6 Hz, CH₃-Ala), 4.50–4.60 (6H, m, 3 × CHO-Oxd + 3 × CHN-Oxd), 5.10 (1H, d, *J* = 6.8 Hz, CH-Ala), 5.36 (2H, s), 5.56 (1H, d, *J* = 6 Hz), 5.59 (1H, d, *J* = 7.6 Hz), 7.41–7.47 (2H, m), 7.28–7.40 (3H, m), 8.37 (1H, s, CH-triazole), 8.40 (1H, s, CH-triazole), 8.59 (1H, s, CH-triazole), 8.59, 114, s, CH-triazole), 8.50, 50.3, 50.5, 50.7, 51.4, 52.1, 54.9, 60.9, 61.8, 69.0, 77.0, 78.4, 81.8, 130.1, 130.7, 131.9, 132.3, 138.2, 141.8, 144.7, 154.6, 159.0, 162.9, 175.6. Anal. Calcd for C₄₅H₅₅N₁₅O₁₅: C, 51.67; H, 5.30; N, 20.09. Found: C, \$1.66; H, \$5.27; N, 20.05.

Boc-(*L*-Ala-L-Oxd-Tri-CO)₃-OH (16). For the synthetic procedure from 15 (0.04 mmol, 42 mg), see the preparation of 6 given above (99% yield, 0.39 mmol, 38 mg): mp = 155 °C; $[\alpha]^{20}_{D} = -23.2$ (*c* = 0.50, CH₃OH); IR (nujol): ν 3335, 1748, 1735, 1684, 1653, 1617, 1577, 1559, 1507 cm⁻¹; ¹H NMR (CD₃OD, 400 MHz) δ 1.13–1.67 (27H, m, 3 × CH₃ Ala + 3 × CH₃ Oxd + *t*-Bu), 3.92–4.01 (3H, m, 3 × CHN), 4.54–4.68 (10H, m, 3 × CHO + 3 × CH₂CH + CH-Ala), 4.95–5.05 (2H, m, 2 × CH-Ala), 8.68 (1H, s, CH-triazole), 8.74 (1H, s, CH-triazole), 8.77 (1H, s, CH-triazole); ¹³C NMR (CD₃OD, 100 MHz) δ 16.1, 19.0, 29.3, 36.0, 37.6, 51.5, 52.2, 52.3, 58.4, 58.5, 75.9, 126.5, 127.1, 142.2, 159.3, 160.6, 172.9. Anal. Calcd for C₃₈H₄₉N₁₅O₁₅: C, 47.75; H, 5.17; N, 21.98. Found: C, 47.79; H, 5.19; N, 21.96.

ECD Spectroscopy. Measurements were performed in a cylindrical, fused quartz cell (1 mm path length). Each pseudopeptide was dissolved in MeOH/H₂O 8:2 v/v at 0.1–1 mM concentration. Metal-binding experiments were performed by adding aliquots of a 65 mM CuSO₄ solution in MeOH/H₂O 8:2 v/v. Ligand competition experiments were performed by adding aliquots of a 0.1 M EDTA solution in H₂O.

UV–vis Spectroscopy. Electronic absorption spectra in the absence and presence of Cu²⁺ were recorded at RT on a doublebeam spectrophotometer, in quartz cells (1 cm-path length). Metal binding experiments were performed by adding aliquots of a 65 mM CuSO₄ solution in MeOH/H₂O 8:2 v/v 0.696 mM pseudopeptides solutions in MeOH/H₂O 8:2 v/v. The dissociation constant (K_D) of the ligand–metal complexes was determined by complexometric titration according to the protocol described by Ferrari et al.²⁸

X-ray Crystallography. Structural investigations by single-crystal X-ray crystallography on 5 and 7.0.5C₆H₆ were carried out on a diffractometer equipped with a Mo K α generator, area detector, and Kryo-Flex liquid-nitrogen cryostat. Crystals were grown from benzene (5) or from benzene/methyl tert-butyl ether 1:1 v/v (7) via vapor diffusion over a Nujol trap. The very long needles so obtained were cut to a proper size using a razor blade, fixed on the tip of a glass capillary with vacuum grease and transferred to the cold nitrogen stream for data collection at 150(2) K. Evaluation of the crystal quality and determination of unit cell parameters were based on 60 preliminary frames collected using ω scans. Data collection was based on ω or $\dot{\phi}$ scans (0.4°) and was extended up to $2\theta = 54^{\circ}$. Data integration, reduction, and correction for absorption were based on standard methods. The structures were solved by direct-methods program SIR92²⁹ and refined on F_0^2 by SHELXL-97³⁰ using the WINGX v2013.3 package.³¹ All non-hydrogen atoms were refined anisotropically (unless otherwise stated), while hydrogen atoms were treated isotropically. In 5, an idealized geometry was assigned to methyl hydrogens, but the torsion angle was allowed to refine freely (AFIX 137 instruction). Their displacement parameter was constrained to be 1.5 times larger than the U_{iso} of the parent C atom. Phenyl hydrogen atoms were subject to unconstrained refinement, while methylene hydrogens on C1 were forced to have the same U value. The remaining H atoms were treated with a common displacement parameter. The absolute structure of 5 could not be reliably determined with Mo K α radiation and was chosen to give S configuration at C12, C13, and C17.

In 7.0.5C₆H₆, the terminal phenyl ring was found disordered over two positions with 0.70:0.30 occupancies. The two components were restrained to have same geometry and to approach mm2 symmetry and were refined isotropically with identical U values for corresponding atoms. An idealized geometry was assigned to methyl hydrogens, whose displacement parameter was constrained to be 1.5 times larger than the U_{iso} of the parent C atom. Torsion angle was however allowed to refine freely (AFIX 137 instruction), except for Boc methyls (AFIX 33). For the remaining hydrogen atoms two common U values were introduced after inspection of the U_{iso} parameters of the attached C or N atoms. Triazole CH hydrogens were set on the N-C-C bisector but the C-H distance was allowed to refine (AFIX 44) and restrained to be similar within 0.02 Å in the two moieties. The amide N-H distances where also restrained to 0.880(15) Å, while C-H distances for tertiary hydrogens on C17 and C28 where forced to be similar within 0.02 Å. After localization of all atoms in the trimer, the unit cell

contained 254 Å³ of potentially solvent-accessible void mainly centered at (0.5, 0.0, 0.0). Electron density residuals (<0.8 e Å³) arranged in an approximately planar fashion were fitted to a half-occupancy benzene molecule, as required for reasonable intermolecular contacts, affording final indices wR2 = 0.1504 (on all data) and R1 = 0.0534 (on data with $I > 2\sigma(I)$). Alternatively, data were treated using SQUEEZE routine in PLATON (V-290610) software to remove the contribution from disordered solvent.³² Final refinement on corrected data gave wR2 =0.1348 (on all data) and R1 = 0.0497 (on data with $I > 2\sigma(I)$) with only slight improvement of bond precision (from 0.0051 to 0.0045 Å for C-C bonds). For this reason, the former approach was preferred. The absolute structure of $7.0.5C_6H_6$ could not be reliably determined with Mo K α radiation and was chosen to give the S configuration at C12, C13, C17, C23, C24, and C28. CCDC 997293-997294 contain the supplementary crystallographic data for this paper. These data can be obtained free of charge from the Cambridge Crystallographic Data Centre via www.ccdc.cam.ac.uk/data request/cif.

EPR Spectroscopy. X-band EPR spectra of the Cu²⁺-peptidomimetic complexes were recorded at 77 K. Instrumental settings: frequency 9.492 GHz; microwave power 5.078 mW modulation amplitude 2.072 G; modulation frequency 100 MHz; magnetic field range 2300–4000 G;. time constant 163.8 ms; conversion time 163.8 ms. Aliquots of a 65 mM CuSO₄ solution in MeOH/H₂O 8:2 v/v were added to 1–5 mM pseudopeptides solutions in MeOH/H₂O 8:2 v/v. Ligand competition experiments were performed by adding aliquots of a 0.1 M EDTA solution in H₂O. Spectral simulations were performed with the EPRSim32 software.³³

Molecular Dynamics Simulations. Molecular dynamics simulations were carried out with the AMBER 11.0 suite of programs.³⁴ The peptide is parametrized using AMBER10 force field³⁴ for the L-Ala residues and GAFF (general AMBER force field) force field³⁵ for the L(D)-Oxd-Tri residues (the standard RESP procedure is carried out to assign charges to atoms by Antechamber).³⁶ A linear conformation of the peptide was built and immersed in a solvent box of explicit methanol molecules. Periodic boundary conditions were used. An equilibration protocol consisting of four individual steps was applied, resulting in an unconstrained well-tempered NPT ensemble at target conditions. A Langevin thermostat was used to set a constant temperature at 300 K and 1 atm. Particle Mesh Ewald³⁷ summation was used throughout (cut off radius of 10 Å for the direct space sum). Bonds involving H atoms were constrained using the SHAKE algorithm,³⁸ and a time step of 2 fs was applied in all runs. Overall sampling time for MD production was 1 μ s. Snapshot structures were saved into individual trajectory files every 5000 time steps, that is, every 10 ps of molecular dynamics, for a total of 100000 snapshots. MD simulations were carried out using pmemd.³⁵ VMD was used to visualize the trajectory.³⁹

Two-dimensional free-energy profiles for the peptide in explicit solvent were obtained as a function of PCA1 and PCA2. The program "ptraj"³⁴ in the AMBER package was used in the PCA. The values are given in kcal mol⁻¹. The energy landscape of the peptide is visualized by means of free-energy functions, which are projected as contour lines onto a two-dimensional space formed by the PCA1/PCA2 axes. These coordinates are derived from a principal component analysis.^{25a} The free-energy change associated with the passage between two different states of a system in thermodynamic equilibrium is given by ΔG = -RT (ln p_1/p_2). Here, R is the ideal gas constant, T is the absolute temperature, and p_i is the probability of finding the system in state *i*. The two-dimensional space defined by the PCA1 and PCA2 axes has been divided into a grid and the free energy has been calculated for each bin of the grid on the basis of the previous equation. The whole set of G values was shifted in such a way that the lowest value of the free energy surface corresponds to zero. Thus, the reported ΔG values represent the transfer free energies with respect to the bin that has been set to zero. To obtain the p values, the trajectory at ambient temperature was projected onto the PCA1/PCA2 space, and p corresponds to the number of times the trajectory "visits" a given bin.

Supporting Information

¹H and ¹³C NMR spectra of compounds 5-16; ROESY spectra of compounds 9 and 15; Tables S1–S4; Figures S1–S6; Supplemental Videos S1 and S2. This material is available free of charge via the Internet at http://pubs.acs.org.

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Notes

The authors declare no competing financial interest.

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REFERENCES

(1) Hill, D. J.; Mio, M. J.; Prince, R. B.; Hughes, T. S.; Moore, J. S. Chem. Rev. 2001, 101, 3893–4011. (b) Hecht, S.; Huc, I. Foldamers: Structure, Properties, and Applications; Wiley-VCH: Weinheim, 2007.

(2) Gellman, S. H. Acc. Chem. Res. 1998, 31, 173-180.

(3) (a) Martinek, T. A.; Fülöp, F. Chem. Soc. Rev. 2012, 41, 687–702.
(b) Cubberley, M. S.; Iverson, B. L. Curr. Opin. Chem. Biol. 2001, 5, 650–653.
(c) Seebach, D.; Beck, A. K.; Bierbaum, D. J. Chem. Bioldivers. 2004, 1, 1111–1239.
(d) Sanford, A.; Yamato, K.; Yang, X. W.; Yuan, L. H.; Han, Y. H.; Gong, B. Eur. J. Biochem. 2004, 271, 1416–1425.
(e) Cheng, R. P. Curr. Opin. Struct. Biol. 2004, 14, 512–520.
(f) Balbo Block, M. A.; Kaiser, C.; Khan, A.; Hecht, S. Top. Curr. Chem. 2005, 245, 89–150.
(g) Goodman, C. M.; Choi, S.; Shandler, S.; DeGrado, W. F. Nat. Chem. Biol. 2007, 3, 252–262.
(h) Bautista, A. D.; Craig, C. J.; Harker, E. A.; Schepartz, A. Curr. Op. Chem. Biol. 2007, 11, 685–692.
(i) Smaldone, R. A.; Moore, J. S. Chem.—Eur. J. 2008, 14, 2650–2657.
(j) Guichard, G.; Huc, I. Chem. Commun. 2011, 47, 5933–5941.

(4) Tomasini, C.; Angelici, G.; Castellucci, N. Eur. J. Org. Chem. 2011, 3648-3669.

(5) (a) Tomasini, C.; Villa, M. *Tetrahedron Lett.* **2001**, *42*, 5211–5214. (b) Bernardi, F.; Garavelli, M.; Scatizzi, M.; Tomasini, C.; Trigari, V.; Crisma, M.; Formaggio, F.; Peggion, C.; Toniolo, C. *Chem.—Eur. J.* **2002**, *8*, 2516–2525.

(6) Tomasini, C.; Luppi, G.; Monari, M. J. Am. Chem. Soc. 2006, 128, 2410–2420.

(7) (a) Sharma, G. V. M; Babu, B. S.; Chatterjee, D.; Ramakrishna, K. V. S.; Kunwar, A. C.; Schramm, P.; Hofmann, H.-J. *J. Org. Chem.* **2009**, 74, 6703–6713.

(8) (a) Hayen, A.; Schmitt, M. A.; Nagassa, N.; Thomson, K. A.; Gellman, S. H. Angew. Chem., Int. Ed. 2004, 43, 505-510. (b) De Pol, S.; Zorn, C.; Klein, C. D.; Zerbe, O.; Reiser, O. Angew. Chem., Int. Ed. 2004, 43, 511-514. (c) Seebach, D.; Jaun, B.; Sebesta, R.; Mathad, R. I.; Flogel, O.; Limbach, M.; Sellner, H.; Cottens, S. Helv. Chim. Acta 2006, 89, 1801-1825. (d) Baldauf, C.; Gunther, R.; Hofmann, H.-J. Biopolymers 2006, 84, 408-413. (e) Sharma, G. V. M; Nagendar, P.; Jayaprakash, P.; Krishna, P. R.; Ramakrishna, K. V. S.; Kunwar, A. C. Angew. Chem., Int. Ed. 2005, 44, 5878-5882. (f) Choi, S. H.; Guzei, I. A.; Gellman, S. H. J. Am. Chem. Soc. 2008, 129, 13780-13781. (g) Prabhakaran, P.; Kale, S. S.; Puranik, V. G.; Rajamohanan, P. R.; Chetina, O.; Howard, J. A. K.; Hofmann, H.-J.; Sanjayan, G. J. J. Am. Chem. Soc. 2008, 130, 17743-17754. (h) Celis, S.; Gorrea, E.; Nolis, P.; Illa, O.; Ortuño, R. M. Org. Biomol. Chem. 2012, 10, 861-868. (i) Pilsl, L. K. A.; Reiser, O. Amino Acids 2011, 41, 709-718.

(9) (a) Price, J. L.; Horne, W. S.; Gellman, S. H. J. Am. Chem. Soc.
2007, 129, 6376–6377. (b) Seebach, D.; Gardiner, J. Acc. Chem. Res.
2008, 41, 1366–1375. (c) David, R.; Gunther, R.; Baumann, L.;

Luhmann, T.; Seebach, D.; Hofmann, H.-J; Beck-Sickinger, A. G. J. Am. Chem. Soc. 2008, 130, 15311–15317.

(10) (a) Sharma, G. V. M; Jadhav, V. B.; Ramakrishna, K. V. S.; Jayaprakash, P.; Narsimulu, K.; Subash, V.; Kunwar, A. C. J. Am. Chem. Soc. 2006, 128, 14657–14668. (b) Baldauf, C.; Gunther, R.; Hofmann, H.-J. J. Org. Chem. 2006, 71, 1200–1208. (c) Chatterjee, S.; Vasudev, P. G.; Ragothama, S.; Shamala, N.; Balaram, P. Biopolymers 2008, 90, 759–771. (d) Vasudev, P. G.; Chatterjee, S.; Ananda, K.; Shamala, N.; Balaram, P. Angew. Chem., Int. Ed. 2008, 47, 6430–6432. (e) Sharma, G. V. M; Babu, B. S.; Ramakrishna, K. V. S.; Nagendar, P.; Kunwar, A. C.; Schramm, P.; Baldauf, C.; Hofmann, H.-J. Chem.—Eur. J. 2009, 15, 5552–5566. (f) Saludes, J. P.; Ames, J. B.; Gervay-Hague, J. J. Am. Chem. Soc. 2009, 131, 5495–5505.

(11) Castellucci, N.; Tomasini, C. Eur. J. Org. Chem. 2013, 3567–3573.

(12) Angelo, N. G.; Arora, P. S. J. Org. Chem. 2007, 72, 7963–7967.
(13) Brik, A.; Alexandratos, J.; Lin, Y.-C.; Elder, J. H.; Olson, A. J.; Wlodawer, A.; Goodsell, D. S.; Wong, C.-H. ChemBioChem 2005, 6, 1167–1169.

(14) (a) Wu, C.-F.; Li, Z.-M.; Xu, X.-N.; Zhao, Z.-X.; Zhao, X.;
Wang, R.-X.; Li, Z.-T. Chem.—Eur. J. 2014, 20, 1418–1426.
(b) Angelo, N. G.; Arora, P. S. J. Org. Chem. 2007, 72, 7963–7967.
(c) You, L.-Y.; Chen, S.-G.; Zhao, X.; Liu, Y.; Lan, W.-X.; Zhang, Y.;
Lu, H.-J.; Cao, C.-Y.; Li, Z.-T. Angew. Chem., Int. Ed. 2012, 51, 1657–1661.
(d) Juwarker, H.; Lenhardt, J. M.; Pham, D. M.; Craig, S. L. Angew. Chem., Int. Ed. 2008, 47, 3740–3743.
(e) Angelo, N. G.; Arora, P. S. J. Am. Chem. Soc. 2005, 127, 17134–17135.

(15) Yuan-Yuan, Liu J. Coord. Chem. 2007, 60, 2597–2605.

(16) Lozano-Vila, A. M.; Luna-Giles, F.; Viñuelas-Zahínos, E.; Cumbrera, F. L.; Ortiz, A. L.; Barros-García, F. J.; Rodríguez, A. B. *Inorg. Chim. Acta* **2011**, 365, 282–289.

(17) Nicoll, A. J.; Miller, D. J.; Futterer, K.; Ravelli, R.; Allemann, R. K. J. Am. Chem. Soc. **2006**, 128, 9187–9193.

(18) (a) Tornoe, C. W.; Christensen, C.; Meldal, M. J. Org. Chem. 2002, 67, 3057–3064. (b) Zhang, L.; Chen, X. G.; Xue, P.; Sun, H. H. Y.; Williams, I. D.; Sharpless, K. B.; Fokin, V. V.; Jia, G. C. J. Am. Chem. Soc. 2005, 127, 15998–15999.

(19) Lemen, G. S.; Wolfe, J. P. Org. Lett. 2010, 12, 2322-2325.

(20) Hu, T.-S.; Tannert, R.; Arndt, H.-D.; Waldmann, H. Chem. Commun. 2007, 38, 3942-3944.

(21) (a) Belvisi, L.; Gennari, C.; Mielgo, A.; Potenza, D.; Scolastico, C. Eur. J. Org. Chem. **1999**, 389–400. (b) Yang, J.; Gellman, S. H. J. Am. Chem. Soc. **1998**, 120, 9090–9091. (c) Jones, I. G.; Jones, W.; North, M. J. Org. Chem. **1998**, 63, 1505–1513. (d) Toniolo, C.; Benedetti, E. CRC Crit. Rev. Biochem. **1980**, 9, 1–44. (e) Imperiali, B.; Moats, R. A.; Fisher, S. L.; Prins, T. J. J. Am. Chem. Soc. **1992**, 114, 3182–3188.

(22) Pokorski, J. K.; Miller Jenkins, L. M.; Feng, H.; Durell, S. R.; Bai, Y.; Appella, D. H. Org. Lett. 2007, 9, 2381–2388.

(23) Toniolo, C.; Formaggio, F.; Tonion, S.; Broxterman, Q. B.; Kaptein, B.; Huang, R.; Setnicka, V.; Keiderling, T. A.; McColl, I. H.; Hecht, L.; Barron, L. D. *Biopolymers* **2004**, *75*, 32–45.

(24) Farrugia, L. J. J. Appl. Crystallogr. 1997, 30, 565-565.

(25) (a) Daidone, I.; Amadei, A. WIREs Comput. Mol. Sci. 2012, 2,

762–770. (b) Fermani, S.; Trivelli, X.; Sparla, F.; Thumiger, A.; Calvaresi, M.; Marri, L.; Falini, G.; Zerbetto, F.; Trost, P. *J. Biol. Chem.* **2012**, *287*, 21372–21383.

(26) (a) Sakaguchi, U.; Addison, A. W. J. Chem. Soc., Dalton Trans. 1978, 601–608. (b) Romanowski, S. M.; Tormena, F.; dos Santos, V.

A.; Hermann, M.; Mangrich, A. S. J. Braz. Chem. Soc. 2004, 897–903. (27) Hathaway, B. J. J. Chem. Soc., Dalton Trans. 1971, 1196–1199.

(28) Ferrari, R. P.; Ghibaudi, E. M.; Traversa, S.; Laurenti, E.; De Gioia, L.; Salmona, M. J. Inorg. Biochem. 1997, 68, 17-26.

(29) Altomare, A.; Cascarano, G.; Giacovazzo, C.; Guagliardi, A. J. Appl. Crystallogr. **1993**, 26, 343-350.

(30) Sheldrick, G. M. Acta Crystallogr. 2008, A64, 112-122.

(31) Farrugia, L. J. J. Appl. Crystallogr. 2012, 45, 849-854.

(32) (a) Spek, A. L. J. Appl. Crystallogr. 2003, 36, 7–13. (b) Spek, A. L. Acta Crystallogr. 2009, D65, 148–155.

(33) Spalek, T.; Pietrzik, P.; Sojca, J. J. Chem. Ifn. Model. 2005, 45, 18–29.

(34) Case, D. A.; Darden, T. A.; Cheatham, T. E., III; Simmerling, C. L.; Wang, J.; Duke, R. E.; Luo, R.; Walker, R. C.; Zhang, W.; Merz, K. M.; Roberts, B.; Wang, B.; Hayik, S.; Roitberg, A.; Seabra, G.; I. Kolossváry, Wong, K. F.; Paesani, F.; Vanicek, J.; Liu, J.; Wu, X.; Brozell, S. R.; Steinbrecher, T.; Gohlke, H.; Cai, Q.; Ye, X.; Wang, J.; Hsieh, M.-J.; Cui, G.; Roe, D.R.; Mathews, D. H.; Seetin, M. G.; Sagui, C.; Babin, V.; Luchko, T.; Gusarov, S.; Kovalenko, A.; Kollman, P.A. AMBER 11, University of California, San Francisco, 2010.

(35) Wang, J.; Wolf, R. M.; Caldwell, J. W.; Kollman, P. A.; Case, D. A. J. Comput. Chem. 2004, 25, 1157–1174.

(36) Wang, J.; Wang, W.; Kollman, P. A.; Case, D. A. J. Mol. Graph. Model. 2006, 25, 247260.

(37) Darden, T.; York, D.; Pedersen, L. J. Chem. Phys. 1993, 98, 10089-10092.

(38) Ryckaert, J.-P.; Ciccotti, G.; Berendsen, H. J. C. J. Comput. Phys. 1977, 23, 327–341.

(39) Humphrey, W.; Dalke, A.; Schulten, K. J. Mol. Graph. 1996, 14, 33–38.