



# Stereoselective synthesis of conformationally constrained reverse turn dipeptide mimetics

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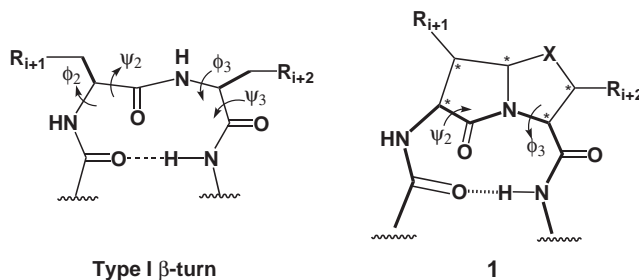
**Abstract**—Peptide side chains play critical roles in the event of molecular recognition. In order to study the bioactive conformation of parent peptides, a concise and straightforward five-step synthesis of [5.5]-bicyclic reverse turn dipeptide mimetic scaffolds with side chain functionality at the  $i+1$  and  $i+2$  positions has been developed. In the bicyclic structure, two dihedral angles ( $\psi_2$  and  $\phi_3$ ) are greatly restricted. © 2000 Elsevier Science Ltd. All rights reserved.

Our knowledge of how receptor–ligand interactions are manifested in biological changes often is very incomplete. A central goal in peptide and protein research is the development of systematic, predictive approaches to the design of peptidomimetics with specific conformational and topographical properties in order to obtain insights into the bioactive conformation of the native peptide on interaction with its receptor. Important peptidic mediators of biological information transduction such as hormones and neurotransmitters have great potential for medical applications, but often have inherent drawbacks such as high degrees of flexibility, biodegradability, and lack of receptor selectivity, which can complicate their use as drugs. Conformational constraints play an important role in rational design of peptides and peptidomimetics that can overcome these problems.<sup>1</sup>

The ‘secondary structure approach’ to de novo design of peptidomimetics is guided by the simple elegance

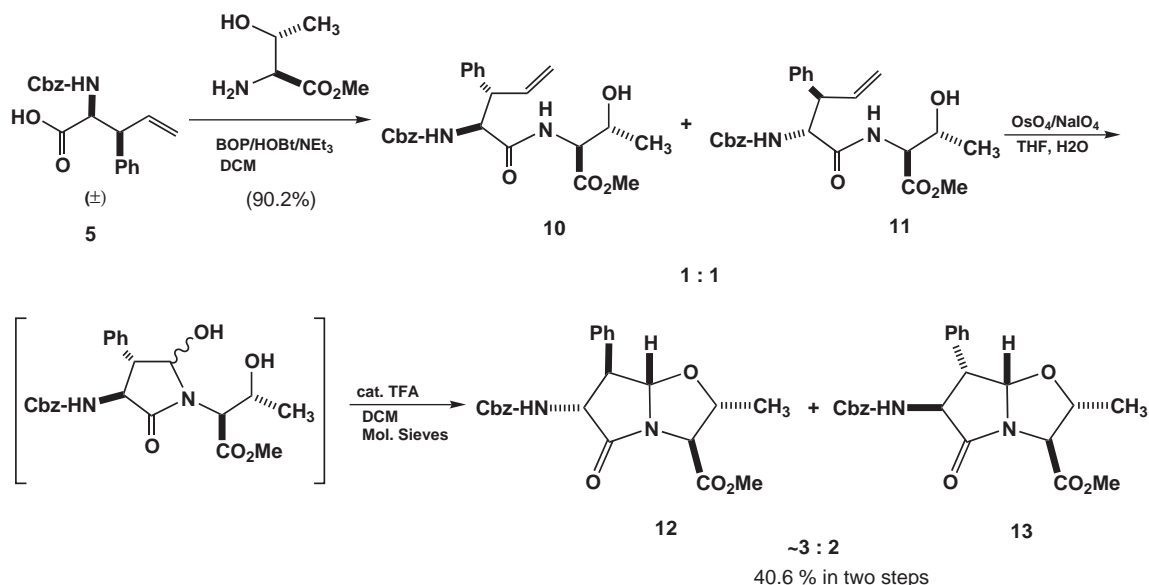
which nature has employed in the molecular architecture of proteins.<sup>2</sup> Of the three major types of secondary structural motifs ( $\alpha$ -helices,  $\beta$ -sheets and reverse turns), reverse turns offer the significant synthetic advantage that they are relatively compact, and of such a size that, in principle, they can be more readily mimicked by conformational constraint, or by use of more rigid small organic molecules.<sup>3</sup>

Though quite a few successes have been reported in obtaining mimetics which can force or stabilize  $\beta$ -turns,<sup>4</sup> very little success has been obtained in incorporating such mimetics into the agonist active site of peptide hormone or neurotransmitter ligands, due to the lack of appropriately positioned side chain groups. In the event of molecular recognition, the peptide backbone serves as a scaffold for the key side chain groups involved in the interaction. The side chain moieties involved directly in the binding are critical for the interaction. Their 3D architecture (topography) and



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Scheme 2.

and **13**. After separation by flash chromatography (EtOAc/hexanes 1:4 to 1:3), the stereochemistry of **12** and **13** was assigned by 1D transient NOE experiments.

In conclusion, a concise and straightforward five-step synthesis of [5.5]-bicyclic reverse turn dipeptide mimetic scaffolds with side chain functionality at the *i*+1 and *i*+2 positions has been developed. In the bicyclic structure, two dihedral angles ( $\psi_2$  and  $\phi_3$ ) are greatly restricted. Further development of the synthesis will enable us to prepare various types of reverse turns with different backbone geometry and side chain topography. Incorporating these conformationally and topographically constrained scaffolds into peptides will help us to understand the bioactive conformation of the parent peptides. Due to its convergent nature, this synthesis also has the potential to be applied to both solid phase chemistry and combinatorial chemistry, which is under investigation.

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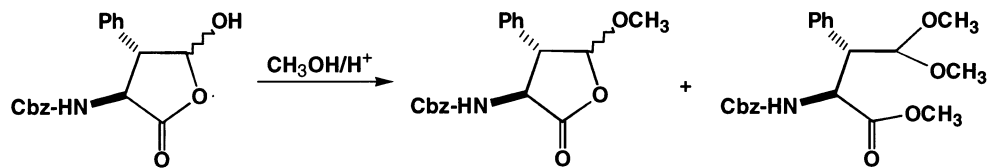
### References

- (a) Hruby, V. J. *Life Sci.* **1982**, *31*, 189–199. (b) Hruby, V. J.; Al-Obeidi, F.; Kazmierski, W. *Biochem. J.* **1990**, *268*, 249–262.
- Nakanishi, H.; Kahn, M. In *Bioorganic Chemistry: Peptides and Proteins*; Hecht, S. M., Ed.; Oxford University Press: London, 1998; pp. 395–419.
- Ripka, W. C.; De Lucca, G. V.; Bach, II, A. C.; Pottorf, R. S.; Blaney, J. M. *Tetrahedron* **1993**, *49*, 3593–3608.
- (a) Kahn, M. Ed. *Tetrahedron* **1993**, *49*, Symposia 50, 3433–3677. (b) Gillespie, P.; Cicariello, J.; Olson, G. L. *Biopolymers* **1997**, *43*, 191–217 and references cited therein. (c) Hanessian, S.; McNaughton-Smith, G.; Lombart, H.-G.; Lubell, W. D. *Tetrahedron* **1997**, *53*, 12789–12854 and references cited therein. (d) Fink, B. E.; Kym, P. R.; Katzenellenbogen, J. A. *J. Am. Chem. Soc.* **1998**, *120*, 4334–4344. (e) Hirschmann, R.; Hynes, J.; Cichy-Knight, M. A.; van Rijn, R. D.; Sprengeler, P. A.; Spoors, P. G.; Shakespeare, W. C.; Pietranico-Cole, S.; Barbosa, J.; Liu, J.; Yao, W.; Rohrer, S.; Smith, A. B. III, *J. Med. Chem.* **1998**, *41*, 1382–1391 and references cited therein. (f) Johannesson, P.; Lindeberg, G.; Tong, W.; Gogoll, A.; Karlen, A. Hallberg, A. *J. Med. Chem.* **1999**, *42*, 601–608. (g) Khalil, E. M.; Ojala, W. H.; Pradhan, A.; Nair, V. D.; Gleason, W. B.; Mishra, R. K.; Johnson, R. L. *J. Med. Chem.* **1999**, *42*, 628–637. (h) Eguchi, M.; Lee, M. S.; Nakanishi, H.; Stasiak, M.; Lovell, S.; Kahn, M. *J. Am. Chem. Soc.* **1999**, *121*, 12204–12205. (i) Gosselin, F.; Lubell, W. D. *J. Org. Chem.* **2000**, *65*, 2163–2171 and references cited therein.
- Hruby, V. J.; Sharma, S. D.; Toth, K.; Jaw, J. Y.; Al-Obeidi, F.; Sawyer, T. K.; de Lauro Castrucci, A.-M.; Hadley, M. E. *Ann. N. Y. Acad. Sci.* **1993**, *680*, 51–63.
- Hruby, V. J.; Han, G. In *The Melanocortin Receptors*; Cone, R. D., Ed.; Humana Press: Totowa, NJ, 2000; pp. 239–261.
- Hruby, V. J.; Chow, M.-S.; Smith, D. D. *Ann. Rev. Pharmacol. Toxicol.* **1990**, *30*, 501–534.
- Hruby, V. J. *Trends Pharmacol. Sci.* **1987**, *8*, 336–339.
- Shenderovich, M. D.; Kövér, K. E.; Wilke, S.; Collins, N.; Hruby, V. J. *J. Am. Chem. Soc.* **1997**, *119*, 5833–5846.
- Al-Obeidi, F.; O'Connor, S. D.; Job, C.; Hruby, V. J.; Pettitt, B. M. *J. Peptide Res.* **1998**, *51*, 420–431.

11. Neises, B.; Steglich, W. *Angew. Chem., Int. Ed. Engl.* **1978**, 17, 522–523.

12. Kazmaier, U. *Synlett* **1995**, 1138–1140.

13.



1 : 9

14. Khalil, E. M.; Pradhan, A.; Ojala, W. H.; Gleason, W. B.; Mishra, R. K.; Johnson, R. L. *J. Med. Chem.* **1999**, 42, 2977–2987.

15. Qiu, W.; Hruby, V. J., unpublished results.

16. Although syntheses of  $\beta$ -hydroxy amino acids have been

extensively studied (for a review, see Genet, J.-P. *Pure Appl. Chem.* **1996**, 68, 593–596), they have limited commercial availability.