

*Communication*

**Evolutionary principles for generating protein mimetics: Directed assembly of peptide loops on topological templates<sup>★</sup>**

Damiano Banfi, Bhubaneswar Mandal, Manfred Mutter and Luc Patiny<sup>✉</sup>

*Institute of Organic Chemistry, University of Lausanne, BCH-Dorigny,  
CH-1015 Lausanne, Switzerland*

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**A novel methodology for the reversible competitive condensation of peptide loops to chemoreactive topological templates is presented.**

The use of topological templates proves to be a versatile concept in protein design [1] and mimicry [2, 3]. By separating the structural and functional part of a protein receptor, the attachment of ligand binding peptide loops to regioselectively addressable template molecules leads to protein mimetics (template-assembled synthetic peptides, Tasp) exhibiting essential features of native receptors [4].

Here, we present a novel methodology for the reversible competitive condensation of peptide loops to chemoreactive topological templates, applying principles of combinatorial chemistry.

**RESULTS AND DISCUSSION**

Based on the design of a metal binding Tasp, a cyclic decapeptide (**T**) as template featuring two aromatic aldehydes for chemoselective ligation and two carboxyl groups as complexing sites was prepared by a convergent strategy (Fig. 1).

N- and C-terminally functionalized linear peptides as prototypes for protein loops (**L1**, **L2**) were reversibly assembled to template **T** *via* imine bond formation (Fig. 2).

To assess the effect of ligand directed template assembly, peptide **L2** featuring two His side chains for metal complexation were com-

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<sup>✉</sup>To whom correspondence should be addressed: Luc.patiny@ico.unil.ch

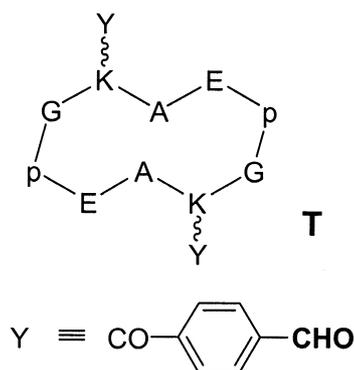


Figure 1

**Loop peptides :**

Xaa : Phe (**L1**); His (**L2**)

DAP = diaminopropionic acid

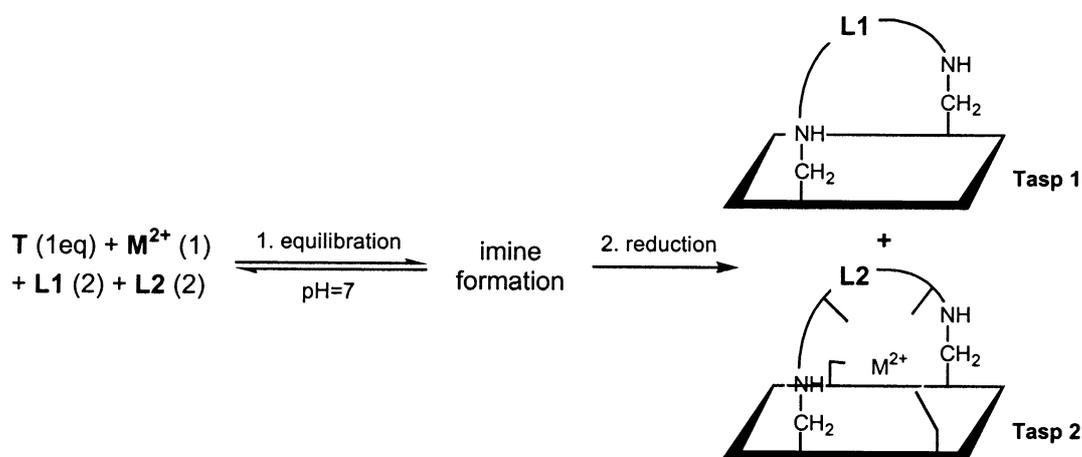


Figure 2

**Table 1.** Yields (%) of Tasp molecules after competitive ligand ( $M^{2+}$ , 1eq) induced assembly of peptide loops L1, L2 (2eq) on a topological template (T, 1eq)

$M^{2+}$	<b>Tasp 1</b>	<b>Tasp 2</b>	Induction factor (IF)
-	47	53	1.00
$\text{Co}^{2+}$	20	80	3.55
$\text{Zn}^{2+}$	30	70	2.07
$\text{Ni}^{2+}$	47	53	1.00

petitively reacted with **L1** (exhibiting no complexing site) in the presence of various metal ions. After equilibration (step 1) the reaction was quenched with  $\text{NaBH}_3\text{CN}$  (step 2) and the resulting Tasp molecules were analysed by HPLC-MS.

In the absence of metal ions, **Tasp 1** and **2** are obtained in about equal amounts pointing to comparable chemical reactivities of **L1** and **L2**. In contrast, a significant preference for

**Tasp 2** is observed in the presence of  $M^{2+}$ , with the metal selectivity (corresponding to the induction factor IF)  $\text{Co} > \text{Zn} \gg \text{Ni}$  (IF = 0).

In conclusion, the elaborated strategy represents a first step in applying evolutionary principles in Tasp design. In particular, the ligand directed assembly of peptide libraries on topological templates opens interesting perspectives in protein mimicry.

## REFERENCES

1. Tuchscherer, G., Grell, D., Mathieu, M. & Mutter, M. (1999) *J. Pept. Res.* **54**, 185–194. [MEDLINE](#)
2. Tuchscherer, G., Scheibler, L., Dumy, P. & Mutter, M. (1998) *Biopolymers* **47**, 63–73. [MEDLINE](#)
3. Sila, U. & Mutter, M. (1995) *J. Mol. Recognit.* **8**, 29–34. [MEDLINE](#)
4. Mutter, M., Dumy, P., Garrouste, P., Lehmann, C., Mathieu, M., Peluso, S., Razaname, A. & Tuchscherer, G. (1996) *Angew. Chem. Int. Ed. Engl* **35**, 1482–1485.