

Ribosomal synthesis of peptide libraries containing cyclic γ -amino acids for drug screening

(¹Graduate School of Science, The University of Tokyo) ○Takashi Miura,¹ Takayuki Katoh,¹ Hiroaki Suga¹

Keywords: *gamma amino acid; unnatural amino acid; macrocyclic peptide; translation; drug discovery*

γ -Amino acids can induce unique secondary structures of peptides, such as C₁₄-helix, C_{12/10}-helix, and C₁₂-turn.^[1] γ -amino acids are found in some natural bioactive peptides and play important roles in their biological activities and proteolytic stability.^[2] Especially, cyclic $\gamma^{2,4}$ -amino acids (c γ AA, Figure) are ideal building blocks that stabilize structures of peptides, and thereby can improve binding affinity to target proteins due to their constrained rigid conformations. Therefore, incorporation of c γ AA into a random peptide library and its application to in vitro selection of active ligands would be powerful approach for developing novel therapeutic peptides.

Recently, our group reported the ribosomal incorporation of c γ AA into nascent peptide chains using a reconstituted translation system assisted by Flexizyme technology, referred to as the Flexible In vitro Translation (FIT) system.^[3] Here we report the ribosomal synthesis of macrocyclic peptide libraries containing two c γ AA, *cis*- and *trans*-3-aminocyclobutane carboxylic acid by means of the FIT system, and its application to an in vitro screening methodology, referred to as the Random nonstandard Peptides Integrated Discovery (RaPID) system.^[4] Consequently, we obtained inhibitor peptides against interferon-gamma receptor 1 (IFNGR1) that showed strong binding affinity and inhibitory activity with low nM *K_D* and IC₅₀ values. Moreover, the peptides containing c γ AA exhibited up to 27-fold longer half-lives in human serum than their alanine-substituted mutants.

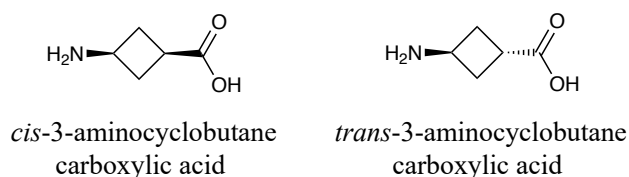


Figure Cyclic γ -amino acids used in this study

[1] Hintermann, T., Gademann, K., Jaun, B., Seebach, D., *Helv. Chim. Acta* **1998**, *81*, 983.

[2] Frackenpohl, J., Arvidsson, P.I., Schreiber, J.V., Seebach, D., *Chembiochem*, **2001**, *2*, 445.

[3] Katoh, T.; Suga, H. *J. Am. Chem. Soc.* **2020**, *142*, 4965.

[4] Yamagishi, Y.; Shoji, I.; Miyagawa, S.; Kawakami, T.; Katoh, T.; Goto, Y.; Suga, H. *Chem. Biol.* **2011**, *18*, 1562.