Efficient Multi-point Interaction between Peptide Catalysts and Amino Acid Esters in a Bilayer Vesicular Membrane for Highly Stereoselective Hydrolysis Reactions

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The ¹H NMR NOESY spectra of the multi-point interaction between *N*-(*N*-benzyloxycarbonyl-L-leucyl)-L-histidine-[(*Z*)-L-Leu-L-His] and methyl *N*-hexanoyl-L-phenylalanate in a *N*,*N*-bisdodecyl-*N*,*N*-dimethylammonium bromide membrane confirms that the membrane-assisted, hydrophobic interaction between the peptide catalyst and enantiomeric substrates enhances the stereoselective hydrolysis of *p*-nitrophenyl *N*-acetyl (or decanoyl)-L (or D)-phenylalanate with (*Z*)-L-Leu-L-His (most effective in di-, tri- or tetra-peptide catalysts with an introduced L-histidyl group).

Although the membrane-assisted stereoselective hydrolysis of amino acid esters has already been performed, mainly with Lhistidyl group-introduced dipeptide catalysts and dialkyl-chain surfactants as a simplified reaction enzymes model,^{1,2} the investigation of the membrane-promoted interaction between an efficient peptide catalyst and an amino-acid substrate in connection with the stereoselective hydrolysis of amino-acid substrates has not been reported. Here we report the determination of the most efficient amino acid sequence in a series of di-, tri- and tetra-peptide catalysts which include nucleophilic Lhistidine and hydrophobic L- (or D)-leucine. We also report on the multi-point interaction between the most efficient peptide catalyst and an amino acid substrate in a bilayer vesicular membrane by means of a 600 MHz ¹H NMR NOESY analysis.

The peptide catalysts 1–5 (see Table 1), the substrates of *p*nitrophenyl *N*-acetyl (or decanoyl)-L- (or D-)-phenylalanate **6** and the surfactant of *N*,*N*-bisdodecyl-*N*,*N*-dimethylammonium bromide 7 were prepared by previously reported methods.¹ The hydrolyses of **6a–c** (10 μ mol dm⁻³) by 1–5 (0.1 mmol dm⁻³) with 7 (1.0 mmol dm⁻³) were carried out in a 3% ν/ν MeCN– Tris buffer (pH 7.68, ionic strength $\mu = 0.15$) at 298 K after sonication of the vesicular membrane (of 7) containing one of 1–5, and the pseduo-first-order hydrolysis rate was followed spectrophotometrically by monitoring the *p*-nitrophenolate anion formation at 400 nm. The catalytic rate constants k_{cat} summarized in Table 1 were obtained by the usual methods.

Among the peptide catalysts tested, the stereoselectivity of the peptide catalysts possessing the L-Leu-L-His unit were

appreciably higher than ones without the L-His-L-Leu unit (1, 2)and 4a) and followed the order $3b \gg 4b > 5a > 5b$. The stereoselectivity of (Z)-L-Leu-L-His 3b was considerably lowered by changing the L-Leu unit into D-Leu (or L-Ala) or by removing the membrane from the reaction system. The catalytic efficiency of 3b was also reduced by changing the substrate from **6b** to C_{10} -L (or D)-Ala-PNP **6c** having a less hydrophobic side-chain. These facts suggest the importance of the membrane-assisted hydrophobic interaction between the side-chains of L-Leu (or L-His) in the peptide catalyst and the enantiomeric substrates for making the reacting positions of the active imidazolyl group of L-His in the catalyst and the susceptible carbonyl group of L (or D)-substrate closer (or further apart), respectively, since the stereoselectivity occured in the catalystsubstrate complex reaction step rather than in the catalystsubstrate binding process.† The 600 MHz ¹H NMR NOESY spectrum (measured by GE NMR OMEGA, mixing time 300 ms) of a D₂O solution (pH 7.0; $\mu = 0.02$) including **3b** (10 mmol dm^{-3}), a substrate analogue of C₆-L-Phe-OMe (methyl *N*-hexanoyl-L-phenylalanate; $10 \text{ mmol } \text{dm}^{-3}$), 7 (20 mmol dm⁻³), and KCl (20 mmol dm⁻³) indicated the multipoint hydrophobic interaction between the L-Leu or L-His sidechain of **3b** and the Ph ring in the side-chain of the substrate. An expected inter-amide hydrogen bonding between the reactants was not observed, probably because of proton-deuterium exchange (Fig. 1).[‡] The same mode of the interaction between 3b and the D-substrate probably makes the reacting groups further apart resulting in the efficient stereoselectivity of 3b. Such a favourable membrane-promoted multi-point interaction

 $k_{cat}/mol^{-1} dm^3 s^{-1}$ 6a 6b Catalyst L L/D D L/D L D (Z)-L-His 1 15 45 1.2 26 1.6 53 17 0.52 (Z-L-His-L-Leu 2 33 87 66 1.3 (Z)-L-Ala-L-His 3a 4.3 7.71.8 41 11 3.7 (Z)-L-Leu-L-His 3b 310 19 16 2590 129 20 (0.99 0.8 1.2)^b 0.007 (0.007) $1.0)^{\circ}$ 108 (Z)-D-Leu-L-His 3c 40 37 1.1 254 2.4 0.91 27 28 0.9 (Z)-L-His-L-Leu 4a 5.2 5.7 149 34 4.4 958 65 15 (Z)-L-Leu-L-His-L-Leu 4b 96 (Z)-L-Leu-L-Leu-L-His 4c 1.8 4.3 268 430 0.62 (Z)-L-Leu-L-His-L-Leu-L-Leu 5a 30 4.4 386 35 11 6.8 (Z)-L-Leu-L-Leu-L-His-L-Leu 5b 28 1.8 291 263 1.1 16

Table 1 Stereoselective hydrolyses of p-nitrophenyl N-acetyl (or decanoyl)-L(or D)-phenylalanate [6a (or 6b)] by peptide catalysts in the membrane of N,N-bisdodecyl-N,N-dimethylammonium bromide 7^a

^{*a*} [Catalyst] = 0.10 mmol dm⁻³, [**6a**-**b**] = 10.0 μ mol dm⁻³, and [**7**] = 1.0 mmol dm⁻³ in 3% ν/ν MeCN-Tris buffer (pH 7.68, μ = 0.15) at 298 K. Experimental errors were within 5%. ^{*b*} Obtained with 3 vol% MeOH instead of 7. ^{*c*} k_{uncat} values obtained without **3b**.

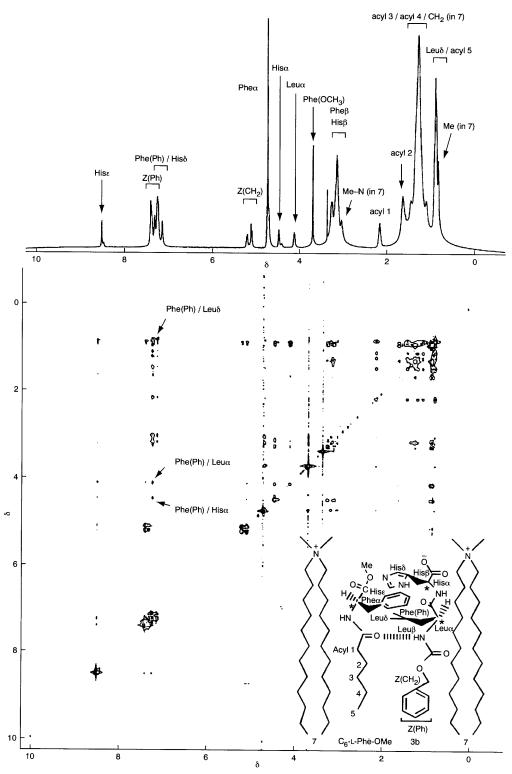


Fig. 1 600 MHz ¹H NMR NOESY spectrum of the interaction between **3b** (10 mmol dm⁻³) and C₆-L-Phe-OMe (10 mmol dm⁻³) in the membrane of **7** in D_2O (pH 7.0) including 20 mmol dm⁻³ KCI ($\mu = 0.02$)

cannot be established by using peptide frameworks other than **3b**.

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Footnotes

 \dagger The enantiomeric ratio (L/D) of the catalyst-substrate (**3b-6b**) binding constant and its reaction rate were calculated to be 1.6 and 8.2, respectively.

References

from stock MeOH solutions.

 K. Ohkubo and S. Miyake, J. Chem. Soc., Perkin Trans. 2, 1987, 995.
Y. Ihara, S. Akasaka, K. Igata, Y. Matsumoto and R. Ueoka, J. Chem. Soc., Perkin Trans. 2, 1991, 543 and references cited therein.

‡ Resonance of the surfactant 7 which was not deuterated was not observed

in the mixing time of 300 ms, presumably because the lateral motion of 7. MeOH (single peak at δ 3.35) which originated not from C₆-L-Phe-OMe but