SYNTHESIS OF TRIPEPTIDES CONTAINING HETEROCYCLIC 
α-AMINO ACIDS BY USING HETEROSPIROCYCLIC 3-AMINO-2H-AZIRINES

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Dedicated to Professor Dr. Kiyoshi Tomioka on the occasion of his 70th birthday

Abstract – By using the ‘azirine/oxazolone method’, di- and tripeptides containing six-membered heterocyclic 4-amino-4-carboxylic acids with a piperidine, tetrahydropyran or tetrahydrothiopyran ring have been synthesized. It has been shown that the corresponding heterospiroyclic 3-(N-methyl-N-phenylamino)-2H-azirines are suitable synthons for these heterocyclic α-amino acids. As expected, the presence of these α,α-disubstituted α-amino acids stabilizes β-turn conformations in the prepared tripeptides of type Z-Phe-Xaa-Val-OR.

INTRODUCTION
The continuing interest in the biological and medicinal chemistry of α,α-disubstituted α-amino acids (Cα-tetrasubstituted α-amino acids) is mainly due to their influence on the conformation of peptides, i.e. their helix-inducing potential.2 In recent years, the enantioselective synthesis of chiral representatives was a focus of several research groups.3 Another type which is of interest consists of heterocyclic n-amino-n-carboxylic acids, for example the 4-amino-4-carboxylate derivatives 1 of piperidine, tetrahydropyran,2 and tetrahydrothiopyran5,6 (Figure 1). Although some studies on amino acids of type 1 as structural units in pharmacologically interesting compounds are known,7 the major interest concerns their insertion into peptides and the resulting structural properties.8
Figure 1. Heterocyclic 4-amino-4-carboxylic acids 1 and heterospirocyclic 3-amino-2H-azirines 2 as synths for 1 in peptide synthesis.

In the last few decades, we evolved an alternative method for the introduction of α,α-disubstituted α-amino acids into peptides, the so-called ‘azirine/oxazolone method’.9 The smooth and efficient coupling of a peptide acid 3 with a 2,2-disubstituted 3-amino-2H-azirine of type 4 yields the extended peptide amide 5 (Scheme 1). Subsequent selective hydrolysis via an intermediate 1,3-oxazolone leads to peptide 6, prepared for the next coupling step. This method was also adapted for solid phase synthesis.10

Scheme 1. Coupling of a peptide acid 3 with 3-amino-2H-azirine 4 and selective hydrolysis to give the extended peptide 6.

The ‘azirine/oxazolone method’ has been used for the synthesis of α-aminoisobutyric (Aib)-containing peptides (peptaibols),11 cyclodepsipeptides12 and cyclopeptides.13 In addition to the Aib-synthon 4, other 2,2-disubstituted 3-amino-2H-azirines of type 4 with various substituents at C(2) have been prepared. Among them, the synthons for heterocyclic α-amino acids14 or dipeptides containing a heterocyclic α-amino acid14c,15 are of special interest.

The goal of the present study was the use of the heterospirocyclic 3-amino-2H-azirines 2a–2c14b (Figure 1) as synthons for heterocyclic α-amino acids 1 in the synthesis of tripeptides as useful units for subsequent peptide syntheses.
RESULTS AND DISCUSSION

According to the standard conditions of the ‘azirine coupling’, a solution of equimolar amounts of a heterospirocyclic 3-amino-2H-azirine $2$ and a N-protected proteinogenic $\alpha$-amino acid $3$ (Phe or Val) in acetonitrile was stirred at room temperature for 16 h. After evaporation of the solvent and chromatographic purification ($\text{SiO}_2$, hexane/AcOEt), the dipeptide amide of type $7$ was obtained in 80–85% yield (Scheme 2, Table 1). The latter compound was dissolved in THF and the solution was cooled to 0 °C. Then, a solution of 6M HCl/THF (1:1) was added and the mixture was stirred at room temperature for 4 h leading to the selective hydrolysis of the terminal amide group. In the cases of $7b$–$e$, the dipeptide acids $8b$–$e$ were isolated in 77–94% yield as colorless crystalline materials. On the other hand, the hydrolysis of the N-Boc-protected $7a$ gave, as expected, $8a$ with the deprotected piperidine ring in 62% yield.

\[
\begin{align*}
3a, b + 2a, c & \xrightarrow{\text{THF or MeCN, rt}} 7a, e \\
\text{PyBOP, Et}_3\text{N, CH}_2\text{Cl}_2, \text{rt} & \xrightarrow{6\text{M HCl / THF (1:1), rt}} 8a, e
\end{align*}
\]

Scheme 2. Synthesis of tripeptides $10$ (Z-Phe-Xaa-Val-OMe and Z-Val-Xaa-Phe-OEt; $Z =$ benzylxycarbonyl)

Finally, $8b$–$e$ were coupled with a second proteinogenic amino acid ester $9$ (Val or Phe) using PyBOP as a classical peptide coupling reagent. The desired tripeptides $10a$–$d$ with protected amino and carboxyl groups were obtained in 89–96% yield. The structures of all prepared compounds were determined on the basis of their spectroscopic and analytical data. Furthermore, the products were formed without epimerization because the $^1\text{H}$- as well as $^{13}\text{C}$-NMR analyses showed that in each case only one stereoisomer was present.
Table 1. Synthesis of tripeptides 10 via the ‘azirine/oxazolone method’

<table>
<thead>
<tr>
<th></th>
<th>R¹</th>
<th>2</th>
<th>X</th>
<th>7</th>
<th>Yield (%)&lt;sup&gt;a&lt;/sup&gt;</th>
<th>8</th>
<th>Yield (%)&lt;sup&gt;a&lt;/sup&gt;</th>
<th>9</th>
<th>R²</th>
<th>R</th>
<th>10</th>
<th>Yield (%)&lt;sup&gt;a&lt;/sup&gt;</th>
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<tbody>
<tr>
<td>a</td>
<td>Bn</td>
<td>a</td>
<td>NBoc</td>
<td>a</td>
<td>82</td>
<td>a&lt;sup&gt;b&lt;/sup&gt;</td>
<td>62</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td></td>
</tr>
<tr>
<td>a</td>
<td>Bn</td>
<td>b</td>
<td>O</td>
<td>b</td>
<td>80</td>
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<td>iPr</td>
<td>b</td>
<td>O</td>
<td>c</td>
<td>82</td>
<td>c</td>
<td>77</td>
<td>b</td>
<td>Bn</td>
<td>Et</td>
<td>b</td>
<td>92</td>
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<tr>
<td>a</td>
<td>Bn</td>
<td>c</td>
<td>S</td>
<td>d</td>
<td>85</td>
<td>d</td>
<td>94</td>
<td>a</td>
<td>iPr</td>
<td>Me</td>
<td>c</td>
<td>94</td>
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<tr>
<td>b</td>
<td>iPr</td>
<td>c</td>
<td>S</td>
<td>e</td>
<td>82</td>
<td>e</td>
<td>80</td>
<td>b</td>
<td>Bn</td>
<td>Et</td>
<td>d</td>
<td>89</td>
</tr>
</tbody>
</table>

<sup>a</sup> Yield of isolated product.  
<sup>b</sup> The dipeptide acid with the deprotected piperidine ring (X = NH) was obtained.

The structures of 10a and 10c were established by single crystal X-ray analysis (Figure 2), and for 10c, the S,S-configuration was confirmed by the refinement of the absolute structure parameter. In the crystal, both tripeptides exist in a β-turn conformation with an intramolecular hydrogen bond between N(4)–H of valine and O(11) of the benzyloxy carbonyl (Z) group (graph set motif<sup>17</sup> S(10); Table 2). The other two NH groups, N(7)–H and N(10)–H, form intermolecular hydrogen bonds with the carbonyl O-atoms O(5) and O(2), respectively, in different neighboring molecules (graph set motifs<sup>17</sup> C(5) and C(11), resp.).

Figure 2. ORTEP plots<sup>16</sup> of the molecular structures of the tripeptides a) Z-Phe-Thp-Val-OMe (10a) and b) Z-Phe-Tht-Val-OMe (10c). Displacement ellipsoids are drawn at the 50% probability level.
Table 2. Intramolecular hydrogen bonds in the tripeptides 10a, 10c and 12a

<table>
<thead>
<tr>
<th>Peptide</th>
<th>H⋯⋯O [Å]</th>
<th>N⋯⋯O [Å]</th>
<th>N–H⋯⋯O [°]</th>
</tr>
</thead>
<tbody>
<tr>
<td>10a</td>
<td>2.27(3)</td>
<td>3.096(3)</td>
<td>158(3)</td>
</tr>
<tr>
<td>10c</td>
<td>2.44(6)</td>
<td>3.154(5)</td>
<td>148(5)</td>
</tr>
<tr>
<td>12a</td>
<td>2.37(3)</td>
<td>3.169(3)</td>
<td>148(2)</td>
</tr>
</tbody>
</table>

The conformations of both tripeptides 10a and 10c are very similar, as shown by the similarity of the torsion angles of the peptide backbone (Figure 3, Table 3). But there is a significant difference between these conformations and those of the previously reported tripeptides Z-Aib-Xaa-Aib-N(Ph)Me₁⁴b which form β-turns of type III with the characteristic torsion angles for Aibᵢ₊₁ and Xaaᵢ₊₂. Whereas the ϕᵢ₊₂/ψᵢ₊₂ values of the heterocyclic amino acids in 10a and 10c are also close to those of a β-turn of type III (+60°/+30°), characteristic for α,α-disubstituted α-amino acids, ϕᵢ₊₁/ψᵢ₊₁ for the Phe residue are indicative of a β-turn of type II (–60°/+120°).

Figure 3. Numbering of the atoms (arbitrary) and torsion angles of the peptide backbone of 10a and 10c

Table 3. Torsion angles [°] within the backbone of the tripeptides 10a, 10c and 12a

<table>
<thead>
<tr>
<th></th>
<th>ωᵢ</th>
<th>φᵢ₊₁</th>
<th>ψᵢ₊₁</th>
<th>ωᵢ₊₁</th>
<th>φᵢ₊₂</th>
<th>ψᵢ₊₂</th>
<th>ωᵢ₊₂</th>
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<tbody>
<tr>
<td>10a</td>
<td>174.3(2)</td>
<td>−52.4(3)</td>
<td>122.8(2)</td>
<td>177.1(2)</td>
<td>54.0(3)</td>
<td>36.9(3)</td>
<td>166.3(2)</td>
</tr>
<tr>
<td>10c</td>
<td>177.2(4)</td>
<td>−57.4(5)</td>
<td>122.3(4)</td>
<td>178.5(3)</td>
<td>51.8(4)</td>
<td>40.4(4)</td>
<td>164.1(3)</td>
</tr>
<tr>
<td>12a</td>
<td>175.7(2)</td>
<td>−51.9(2)</td>
<td>124.1(2)</td>
<td>175.2(2)</td>
<td>52.0(2)</td>
<td>40.3(2)</td>
<td>162.5(2)</td>
</tr>
</tbody>
</table>

There is a strong indication that the tripeptide derivatives 10a–d also exist as β-turns in solution, similar to the previously reported tripeptides Z-Aib-Xaa-Aib-N(Ph)Me containing only α,α-disubstituted α-amino acids.¹⁴b The three NH signals in the ¹H-NMR spectra in CDCl₃ of all four components could be assigned to N(4)H (7.4–7.1 ppm), N(7)H (6.19–6.57 ppm) and N(10)H (5.34–5.49 ppm) on the basis of their multiplicity and 2D-NMR experiments. It is well known that the chemical shifts of NH groups of
peptides involved in intramolecular hydrogen bonds show a small dependence on the solvent polarity, whereas those exposed to the solvent are influenced significantly. The measurements in CDCl$_3$ containing 0–12% (D$_6$)DMSO at ca. 30 °C showed a very small solvent-dependence for N(4)H ($\Delta\delta = 0.1–0.6$ ppm) but a significant one in the cases of N(7)H and N(10)H ($\Delta\delta = 1.14–1.50$ ppm). We explain these results with the presence of an intramolecular hydrogen bond N(4)–H…O(11), i.e. a $\beta$-turn conformation also in solution.

The problem arising in the hydrolysis of the dipeptide amide 7a bearing the acid-labile Boc-protecting group in the piperidine ring was not unexpected. The attempts to reprotect the resulting dipeptide acid 8a ($X = \text{NH}$) with (Boc)$_2$O under standard conditions were unsuccessful. On the other hand, the reaction with Z-chloride led to the N-protected dipeptide acid 11 as a useful building block for peptide synthesis in 39% yield (Scheme 3).

\[\text{8a} \rightarrow \text{Z-Cl, 0.5M NaOH, H$_2$O/Dioxane (1:1)} \rightarrow \text{Z-N} \begin{array}{c} \text{H} \\ \text{O} \end{array} \text{N} \begin{array}{c} \text{H} \\ \text{O} \end{array} \text{H} \begin{array}{c} \text{O} \\ \text{Ph} \end{array} \text{11} \]

**Scheme 3.** Reprotection of the piperidine N-atom with benzyloxycarbonyl chloride (Z–Cl)

With the aim of obtaining tripeptide units useful for further peptide synthesis, the selective deprotections of the terminal carboxylic acid and amino group, respectively, were studied. Treatment of 10a–d with LiOH·H$_2$O in THF/MeOH/H$_2$O (2:1:1) at room temperature for 2 h, acidic workup and crystallization led to the tripeptide acids 12a–d in very good yields (91–99%, Scheme 4). All products were obtained as single diastereoisomers evidenced by a single set of signals in the $^1$H- and $^{13}$C-NMR spectra. Therefore, the saponification proceeded under preservation of the configuration.

The hydrogenolytic deprotection of the N-terminus of the tripeptides 10a,b by treatment with Pd/C in methanol gave the desired products 13a,b in high yields (Scheme 4). On the other hand, the analogous reactions with the S-containing tripeptides 10c,d were unsuccessful, and also under modified conditions, the deprotection could not be achieved. As a consequence, for further peptide synthesis with the Tht-synthon 2c, another N-protecting group has to be chosen.
Scheme 4. Selective deprotection of the carboxylic acid and amino terminus, respectively, of the tripeptide derivatives 10a-d

In the case of 12a, the molecular structure was established by X-ray crystallography (Figure 4). The crystals contain one molecule of water for every peptide molecule. The carboxylic acid group forms an intermolecular hydrogen bond with the O-atom of the water molecule, which in turn forms intermolecular hydrogen bonds with the second amide O-atom and the third amide N-atom of two different peptide molecules. These interactions respectively link the peptide and water molecules in an alternating sequence into extended chains which run parallel to the [001] and [−101] directions. The graph set motifs$^{17}$ describing these chains are $C_2(12)$ and $C_2(13)$, respectively. N(10)–H forms an intermolecular hydrogen bond with the carboxylate O-atom, O(2), of a neighboring peptide molecule to build a chain which runs parallel to the [001] direction with a graph set motif of C(11). N(7)–H forms an intermolecular hydrogen bond with the amide O-atom, O(5), of a different adjacent peptide molecule thereby building a chain which runs parallel to the [100] direction with a graph set motif of C(5). The combination of all the intermolecular interactions links the molecules into two-dimensional sheets, which lie parallel to the (010) plane. Finally, N(4)–H forms an intramolecular hydrogen bond with the carbonyl O(11)-atom that is seven atoms back along the peptide backbone (graph set S(10)) forming a β-turn (Tables 2 and 3). The conformation of 12a in the crystal is virtually identical with that of the corresponding peptide ester 10a.
CONCLUSIONS

The present studies show that the heterospirocyclic 3-(N-methyl-N-phenylamino)-2H-azirines 2a–c are suitable building blocks for the synthesis of peptides containing heterocyclic 4-amino-4-carboxylic acids with a piperidine, tetrahydropyran or tetrahydrothiopyran ring. According to the ‘azirine/oxazolone method’, their coupling with Z-Val-OH or Z-Phe-OH occurs smoothly and in high yields. The selective hydrolysis of the terminal amide group of the dipeptides and further coupling with a proteinogenic amino acid ester can be achieved easily in the cases of the O- and S-containing heterocyclic amino acids. On the other hand, the acid catalyzed hydrolysis of the terminal amide group of the piperidine-containing dipeptide leads to the deprotection of the piperidine ring. Whereas the tripeptide with a tetrahydropyran residue can be selectively deprotected at the carboxylic acid and amino terminus, respectively, the hydrogenolytic deprotection of the amino group of the tripeptide with the sulfur-containing heterocycle is not possible. The goal of further studies is the evaluation of different protecting groups in the synthon 2a for 4-aminopiperidine-4-carboxylate (1a) as well as for the used amino acids in coupling reactions with the tetrahydrothiopyran-containing synthon. Finally it has been shown that the presence of the heterocyclic α-amino acids 1b and 1c in the prepared tripeptides stabilize β-turn conformations as expected for α,α-disubstituted α-amino acids.

EXPERIMENTAL

General remarks. Melting points were determined using a Mettler FP5 apparatus, and they are
uncorrected. Thin layer chromatography (TLC): Merck silica gel 60 F254 plates (0.25 mm); column chromatography (CC): silica gel Merck 60 (0.040–0.063 mm. The IR spectra were recorded on a Perkin-Elmer 297 or Perkin-Elmer 781 spectrophotometer in CDCl3 or in KBr; absorptions in cm⁻¹. The ¹H- and ¹³C-NMR spectra were measured on a Bruker ARX-300, AM-400 or AMX-600 instrument (300/75.4, 400/100.6, and 600/150.9 MHz, resp.) in CDCl3 with TMS as internal standard. Chemical shifts (δ) are given in ppm and coupling constants J in Hz. Mass spectra (MS) were recorded on a Finnigan SSQ-700 (CI, NH₃, 150 eV) or Finnigan TSQ-700 (ESI) instrument. Optical rotations [α]D: Zeiss LEP-A2 polarimeter, in MeOH at 20–22 °C.

Starting materials. The synthesis of tert-butyl 2-[methyl(phenyl)amino]-1,6-diazaspiro[2.5]oct-1-ene-6-carboxylate (2a), N-methyl-N-phenyl-6-oxa-1-azaspiro[2.5]oct-1-en-2-amine (2b), and N-methyl-N-phenyl-6-thia-1-azaspiro[2.5]oct-1-en-2-amine (2c) has been described previously.¹⁴b The amino acid derivatives Z-Phe-OH, Z-Val-OH, H-Phe-OEt, and H-Val-OMe as well as all used reagents were commercially available. Reported yields refer to isolated products.

Abbreviations. AcOEt, ethyl acetate; Boc, tert-butyloxycarbonyl; MeCN, acetonitrile; Phe, phenylalanine; Pip, 4-aminopiperidine-4-carboxylic acid; PyBOP, [(benzotriazol-1-yl)oxy]tripyrrolidinophosphonium hexafluorophosphate; THF, tetrahydrofuran; Thp, 4-aminotetrahydropyran-4-carboxylic acid; Tht, 4-aminotetrahydrothiopyran-4-carboxylic acid; Val, valine; Z, benzyloxycarbonyl.

General procedure for the synthesis of dipeptide amides 7a–e (azirine coupling). To a solution of the corresponding azirine 2 in MeCN or THF at 0 °C, Z-Phe-OH or Z-Val-OH were added and the mixture stirred at rt for 16 h. Then, the solvent was evaporated and the residue was purified by column chromatography (CC).

Z-Phe-Pip(Boc)-N(Ph)Me (7a). The reaction of 500 mg (1.58 mmol) azirine 2a and 473 mg (1.58 mmol) Z-Phe-OH in 5 mL MeCN followed by CC (hexane/AcOEt 2:1) gave 890 mg (82%) of 7a. Colorless solid; mp 126–127 °C. IR (CHCl3): 3420 w, 3060 w, 3020 w, 3000 m, 2980 w, 2930 w, 2860 w, 1685 s, 1645 m, 1595 m, 1495 s, 1455 m, 1430 m, 1390 w, 1370 m, 1285 m, 1260 m, 1250 m, 1145 m, 1130 w, 1075 w, 1050 w, 1030 w, 995 w, 860 w, 700 m. ¹H-NMR (CDCl3): 7.37–7.02 (m, 15 arom. H, 1 NH); 5.45 (d, J = 6.7, 1 NH); 5.08, 5.01 (2d, JAB = 12.1, PhCH2O); 3.95–3.85 (m, HC(2) of Phe); 3.75–3.55 (m, 2 Heq of –CH2NCH2–); 3.16 (s, MeN); 3.02–2.96 (m, 2 Hax of –CH2NCH2–); 2.65–2.25 (m, CH2 of Phe); 2.09–1.83 (m, –CH2CCH2–); 1.43 (s, Me3C). ¹³C-NMR (CDCl3): 171.2, 169.7 (2s, 2 C=O); 156.1, 154.5 (2s, 2 OCON); 144.6, 136.6, 136.0 (3s, 3 arom. C); 129.6, 129.2, 128.9, 128.6, 128.4, 128.1, 127.9, 127.4, 127.2 (9d, 15
Z-Phe-Thp-N(Ph)Me (7b). The reaction of 652 mg (3.015 mmol) azirine 2b and 905 mg (3.023 mmol) Z-Phe-OH in 8 mL MeCN followed by CC (hexane/AcOEt 2:1) gave 1.132 g (82%) of 7b. Colorless crystals; mp 143–144 °C. IR (CHCl₃): 3420 cm⁻¹.

Z-Phe-Thp-N(Ph)Me (7c). The reaction of 641 mg (2.964 mmol) azirine 2c and 752 mg (2.993 mmol) Z-Phe-OH in 8 mL MeCN followed by CC (hexane/AcOEt 2:1) gave 1.243 g (80%) of 7c. Colorless crystals; mp 172–173 °C. IR (CHCl₃): 3420 cm⁻¹.

Z-Val-Tht-N(Ph)Me \((7e)\). The reaction of 403 mg (1.734 mmol) azirine \(2c\) and 438 mg (1.743 mmol) Z-Val-OH in 5 mL THF followed by CC (hexane/AcOEt 2:1) gave 692 g (82%) of \(7e\). White solid; mp 168–169 °C. IR (CHCl\(_3\)): 3410 \(\nu\) (OH). \(\text{H-NMR (CDCl}_3\)): 7.43–7.26 (m, 10 arom. H); 5.50 (s, 1 NH); 5.42 (d, \(J = 8.8, 1\) NH); 5.18, 5.09 (2d, \(J = 12.3\) Hz, PhCH\(_2\)OH); 3.63 (dd, \(J = 8.8, 4.8\) Hz, HC(2) of Val); 3.20 (s, MeN); 2.62–2.50, 2.40–2.35, 2.24–2.20, 2.08–2.02 (4m, Me\(_2\)CH of Val, 4 CH\(_2\)); 0.94, 0.90 (2d, \(J = 6.8\), Me\(_2\)CH of Val). \(\text{C-NMR (CDCl}_3\)): 171.4, 170.4 (2s, 2 C=O); 156.4 (s, OCON); 144.5, 136.2 (2s, 2 arom. C); 129.6, 128.6, 128.3, 128.1, 127.9, 127.5 (6d, 10 arom. CH); 67.2 (t, PhCH\(_2\)OH); 59.8 (d, HC(2) of Val); 59.7 (s, C(4) of Tht); 41.4 (q, MeN); 35.0, 33.8 (2t, –CH\(_2\)C\(_6\)H\(_5\)–); 31.1 (d, Me\(_2\)CH of Val); 23.7, 23.3 (2t, –CH\(_2\)SCH\(_2\)–). CI-MS: 484 (2, [\(M+1\]^+]\(^+\)), 377 (22), 376 (100, [\(M-N(Ph)Me\)]\(^+\)). \(\alpha\) \(\sim\) 24.6 (c 0.977).

**General procedure for the hydrolysis of dipeptide amides \(7a\)–\(e\).** To a solution or suspension of the corresponding dipeptide amide \(7\) in THF at 0 °C, the same volume of 6M HCl was added and the mixture was stirred at rt for 4 h. Then, the mixture was extracted with CH\(_2\)Cl\(_2\), the organic layer was dried with MgSO\(_4\) and the solvent evaporated in vacuo. The residue was purified by CC or crystallization.

**Z-Phe-Pip-OH (8a).** To a suspension of 1.124 g (1.828 mmol) \(7a\) in 6 mL THF at 0 °C was added 6M HCl (6 mL). The organic solvent was evaporated and the aqueous solution was cooled to 0 °C. Then, 5M NaOH was added until pH 13, the mixture was washed with Et\(_2\)O (2x) and neutralized by addition of 1M HCl. The precipitate was filtered and washed with water to give 630 mg (81%) of \(8a\). Colorless powder; mp 240–243 °C. IR (CHCl\(_3\)): 3300, 3220, 3140, 3060, 3020, 2960, 2940, 2800, 1690, 1680, 1665, 1585, 1540, 1505, 1495, 1465, 1450, 1435, 1410, 1390, 1360, 1350, 1320, 1290, 1240, 1185, 1140, 1080, 1055, 1025, 995, 935, 910, 850, 800, 780, 750, 695. \(\text{H-NMR (CDCl}_3\)): 7.43–7.26 (m, 10 arom. H); 5.20 (s, PhCH\(_2\)OH); 4.85–4.70 (m, HC(2) of Phe); 3.70–3.05 (m, –CH\(_2\)NCH\(_2\)–, CH\(_2\) of Phe); 2.80–2.25 (m, –CH\(_2\)C\(_6\)H\(_5\)–). \(\text{C-NMR (CF}_3\text{CO}_2\text{D)}: 176.1, 174.7 (2s, 2 C=O); 158.1 (s, OCON); 134.3, 131.1 (2s, 2 arom. C); 129.3, 128.8, 128.3, 128.1, 127.6, 127.4 (6d, 10 arom. CH); 68.7 (t, PhCH\(_2\)OH); 55.8 (s, C(4) of Pip); 55.3 (d, HC(2) of Phe); 40.7, 37.3 (2t, –CH\(_2\)NCH\(_2\)–, CH\(_2\) of Phe); 28.1 (t, –CH\(_2\)C\(_6\)H\(_5\)–). ESI-MS: 448 (93, [\(M+Na\)]\(^+\)), 426
(100, [M+1]⁺).

Z-Phe-Thp-OH (8b). Treatment of 1.212 g (2.351 mmol) of 7b in 8 mL THF with 6M HCl (6 mL) at 0 °C followed by CC (CH2Cl2/Methanol/AcOH 100:4:1) gave 872 mg (87%) of 8b. Colorless crystals; mp 109 °C. IR (CHCl3): 3300w, 3030m, 2960m, 2860m, 1720m, 1670s, 1540s, 1455m, 1445m, 1390m, 1290m, 1250s, 1145m, 1105m, 1050m, 1030m, 840m, 740m. 1H-NMR (CD3OD): 7.30–7.18 (m, 10 arom. H); 5.01 (s, PhCH2O); 4.49–4.44 (m, HC(2) of Phe); 3.72–3.64, 3.54–3.47, 3.39–3.29, 3.12–3.05, 2.90–2.82 (5m, –CH2OCH2–, CH2 of Phe); 2.10–1.88 (m, –CH2CH2–). 13C-NMR (CD3OD): 176.3, 173.8 (2s, 2 C=O); 158.1 (s, OCON); 138.5, 138.2 (2s, 2 arom. C); 130.5, 129.5, 129.45, 128.7, 127.8 (5d, 10 arom. CH); 67.6, 64.4 (2t, PhCH2O, –CH2CH2–); 57.7 (s, C(4) of Thp); 57.5 (d, HC(2) of Phe); 39.1, 33.4, 33.2 (3t, CH2 of Phe, –CH2CH2–). ESI-MS: 449 ([M+Na]⁺). [α]D –2.4 (c 1.013). Anal. Calcd for C23H28N2O6 (426.47): C 64.78, H 6.14, N 6.57. Found: C 64.54, H 5.99, N 6.79.

Z-Val-Thp-OH (8c). Treatment of 1.089 g (2.329 mmol) of 7c in 6 mL THF with 6M HCl (6 mL) at 0 °C followed by CC (CH2Cl2/Methanol/AcOH 100:4:1) gave 682 mg (77%) of 8c. Colorless crystals; mp 85–86 °C. IR (KBr): 3300w, 3060m, 3030m, 2960m, 2870m, 1710s, 1670s, 1540s, 1470m, 1455m, 1445m, 1430m, 1390m, 1290m, 1250s, 1145m, 1105m, 1030m, 1000w, 840m. 1H-NMR (CD3OD): 7.35–7.26 (m, 5 arom. H); 5.12, 5.07 (2d, JAB = 12.3, PhCH2O); 3.98 (d, J = 7.3, H(C2) of Val); 3.75–3.55 (m, –CH2OCH2–); 2.16–1.93 (m, Me3CH of Val, –CH2CH2–); 0.99, 0.55 (2d, J = 6.8, Me4CH of Val). 13C-NMR (CDCl3): 174.9, 172.6 (2s, 2 C=O); 157.0 (s, OCON); 136.7 (s, 1 arom. C); 128.0, 127.5, 127.3 (3d, 5 arom. CH); 66.2, 63.0, 62.9 (3t, PhCH2O, –CH2OCH2–); 60.4 (d, HC(2) of Val); 56.0 (s, C(4) of Thp); 32.0, 31.6 (2t, –CH2CH2–); 30.5 (d, Me3CH of Val); 18.2, 17.0 (2q, Me2CH). ESI-MS: 779 (100, [2M+Na]⁺), 401 (58, M+Na⁺). [α]D –15.8 (c 0.943). Anal. Calcd for C19H26N2O6 (378.43): C 60.30, H 6.93, N 7.40. Found: C 60.45, H 7.12, N 7.20.

Z-Phe-Tht-OH (8d). Treatment of 822 mg (1.546 mmol) of 7d in 8 mL THF with 6M HCl (8 mL) at 0 °C followed by crystallization from MeOH/Et2O gave 643 mg (94%) of 8d. Colorless crystals; mp 136–137 °C. IR (CHCl3): 3420m, 3320m, 3060m, 3005m, 2960m, 2920m, 2820m, 1715s, 1510s, 1455m, 1445m, 1430m, 1280m, 1240m, 1145w, 1110w, 1080w, 1050w, 910w, 700s. 1H-NMR (CD3OD): 7.42–7.25 (m, 5 arom. H); 5.09 (s, PhCH2O); 4.55 (dd, J = 9.0, 5.9, HC(2) of Phe); 3.22–3.14, 2.98–2.87, 2.78–2.65, 2.57–2.31, 2.25–2.13 (5m, CH2 of Phe; 4 CH3). 13C-NMR (CD3OD): 174.1, 171.3 (2s, 2 C=O); 155.7 (s, OCON); 136.0, 135.7 (2s, 2 arom. C); 128.0, 127.0, 126.5, 126.3, 125.3 (5d, 10 arom. CH); 65.2 (t, PhCH2O); 56.9 (s, C(4) of Tht); 55.1 (d, HC(2) of Phe); 36.6 (t, CH2 of Phe); 31.9, 31.8 (2t, –CH2CH2–); 21.7 (t, –CH2SCH2–). CI-MS: 460 (13, [M+NH4]⁺), 443 (42, [M+H]⁺), 426 (24), 425 (100, [M−OH]⁺). [α]D–7.7 (c 1.004). Anal. Calcd for C29H26N2O6S (442.54): C 62.42, H 5.92, N 6.33, S 7.25. Found: C 62.17, H 6.08, N 6.00, S 6.95.

Z-Val-Tht-OH (8e). Treatment of 867 mg (1.793 mmol) of 7e in 8 mL THF with 6M HCl (8 mL) at 0 °C
followed by crystallization from MeOH/Et₂O gave 565 mg (80%) of 8e. Colorless crystals; mp 135–136 °C. IR (CHCl₃): 3420w, 3320w, 3020w, 3000w, 2960m, 2920m, 2870m, 1715s, 1510s, 1470w, 1455w, 1445w, 1430w, 1390w, 1370w, 1280m, 1090w, 1025w, 910w, 700m. ¹H-NMR (CDCl₃): 7.33–7.28 (m, 5 arom. H); 7.07 (s, 1 NH); 5.92 (d, J = 8.3, 1 NH); 5.10 (s, PhCH₂O); 4.05–4.00 (m, HC(2) of Val); 2.70–2.50, 2.34–2.07 (2m, Me₂CH of Val, 4 CH₂); 0.92, 0.88 (2d, J = 6.9, Me₂CH of Val). ¹³C-NMR (CDCl₃): 175.7, 172.0 (2s, 2 C=O); 157.0 (s, OCON); 136.0 (s, 1 arom. C); 128.4, 128.1, 127.8 (3d, 5 arom. CH); 67.2 (t, PhCH₂O); 60.4 (d, HC(2) of Val); 58.0 (s, OCON); 33.3, 32.8 (2t, −CH₂CCH₂−); 31.5 (d, Me₂CH of Val); 23.3, 22.6 (2t, −CH₂SCH₂−); 19.2, 17.8 (2q, Me₂CH). CI-MS: 412 (40, [M+NH₄⁺]), 396 (21), 395 (100, [M+1⁺]), 377 (44, [M−OH⁺]). [α]D −9.54 (c 1.007).

**General procedure for the coupling of 8b–e with an amino acid ester to give tripeptide esters 10.** To a solution of the corresponding dipeptide 8 in CH₂Cl₂, 1.1 mol equivalent of Val-OMe.HCl (9a) or Phe-OEt.HCl (9b), respectively, 1 equivalent of the coupling reagent PyBOP, and 3 equivalents Et₃N were added at rt. The mixture was stirred at rt for 1–2 h, then, the solvent was evaporated in vacuo, and the residue was purified by CC.

**Z-Phe-Thp-Val-OMe (10a).** The reaction of 708 mg (1.660 mmol) of 8b, 307 mg (1.831 mmol) 9a, 868 mg (1.668 mmol) PyBOP, and 700 μL (5.0 mmol) Et₃N in 10 mL CH₂Cl₂ at rt, followed by CC (Et₂O/AcOEt 2:1) gave 864 mg (96%) of 10a. Colorless crystals; mp 147–148 °C. IR (CHCl₃): 3420m, 3000m, 2960m, 1735s, 1680s, 1510s, 1470m, 1455m, 1390w, 1370m, 1350m, 1300w, 1240m, 1180w, 1155m, 1110m, 1080w, 1050m, 1030m, 1000w, 910w. ¹H-NMR (CDCl₃): 7.41–7.20 (5m, 1 NH); 5.07 (s, PhCH₂O); 4.47–4.39 (m, HC(2) of Phe, HC(2) of Val); 3.73–3.59, 3.37–3.29, 3.21–3.00 (3m, Me₂OCH₂−, CH₂ of Phe); 3.67 (s, MeO); 2.23–1.98, 1.73–1.68 (2m, −CH₂CCH₂−, Me₂CH of Val); 0.92, 0.88 (2d, J = 6.8, Me₂CH of Val). ¹³C-NMR (CDCl₃): 172.5, 172.2, 171.5 (3s, 3 C=O); 156.2 (s, OCON); 136.1, 135.8 (2s, 2 arom. C); 129.1, 128.8, 128.5, 128.2, 127.9, 127.2 (6d, 10 arom. CH); 67.2 (t, PhCH₂O); 63.0, 62.8 (2t, −CH₂OCH₂−); 57.9 (s, C(4) of Thp); 57.4, 56.8 (2d, HC(2) of Phe, HC(2) of Val); 51.9 (s, MeO); 37.7, 33.1, 31.5 (3t, CH₂ of Phe, −CH₂CCH₂−); 30.8 (d, Me₂CH of Val); 19.0, 17.7 (2q, Me₂CH of Val). ESI-MS: 562 ([M+Na⁺]). [α]D −7.9 (c 1.126). Anal. Calcd for C₂₃H₃₁N₅O₇ (539.63): C 64.55, H 6.91, N 7.72. Found: C 64.44, H 6.78, N 7.90.

**Z-Val-Thp-Phe-OEt (10b).** The reaction of 526 mg (1.390 mmol) of 8c, 353 mg (1.537 mmol) 9b, 727 mg (1.397 mmol) PyBOP, and 600 μL (4.3 mmol) Et₃N in 8 mL CH₂Cl₂ at rt, followed by CC (Et₂O/AcOEt 3:1) gave 711 mg (92%) of 10b. Colorless crystals; mp 125–126 °C. IR (CHCl₃): 3420m, 3300m, 3000m, 2970m, 2940m, 2860m, 1730s, 1680s, 1510s, 1470m, 1455m, 1390m, 1350m, 1290m, 1160m, 1110m, 1080w, 1030m, 980w, 910w, 850w, 825w, 700m. ¹H-NMR (CDCl₃): 7.42 (d, J = 6.6, 1 NH); 7.37–7.12 (m,
10 arom. H); 6.57 (s, 1 NH); 5.36 (d, J = 7.6, 1 NH); 5.09 (s, PhCH2O); 4.75 (q, J = 6.8, HC(2) of Phe); 4.08 (q, J = 7.2, MeCH2O); 3.91 (t, J = 7.3, H(C2) of Val); 3.75–3.64, 3.53–3.46 (2m, –CH2OCH2–); 3.15–2.99 (m, CH2 of Phe); 2.19–1.97, 1.83–1.79 (2m, Me2CH of Val, –CH2CCH2–); 1.17 (t, J = 7.2, MeCH2O); 0.94, 0.90 (2d, J = 6.8, Me2CH of Val). 13C-NMR (CDCl3): 172.3, 171.9, 171.3 (3s, 3 C=O); 156.6 (s, OCON); 136.1, 135.8 (2s, 2 arom. C); 129.2, 128.5, 128.3, 128.2, 127.9, 126.8 (6d, 10 arom. CH); 67.2 (t, PhCH2O); 63.1, 61.2 (2t, –CH2OCH2–, MeCH2O); 61.0 (d, HC(2) of Phe); 57.7 (s, C(4) of Tht); 53.5 (d, HC(2) of Val); 37.9, 32.8, 31.9 (3t, CH2 of Phe, –CH2CCH2–); 30.2 (d, Me2CH of Val); 19.4, 17.5 (2q, Me2CH of Val); 13.9 (q, MeCH2O). ESI-MS: 576 (100, [M+Na]+), 554 (21, [M+1]+). [α]D –15.6 (c 0.996). Anal. Calcd for C30H37N3O6S (555.70): C 62.68, H 6.71, N 7.56, S 5.77. Found: C 62.68, H 6.80, N 7.59. Found: C 65.05, H 7.23, N 7.69.

Z-Phe-Tht-Val-OMe (10c). The reaction of 446 mg (1.008 mmol) of 8d, 187 mg (1.115 mmol) 9a, 524 mg (1.008 mmol) PyBOP, and 420 μL (3.0 mmol) Et3N in 2 mL CH2Cl2 at rt, followed by CC (hexane/AcOEt 1:1) gave 525 mg (94%) of 10c. Colorless crystals; mp 160–161 °C. IR (CHCl3): 3420m, 3005m, 2960m, 2930m, 2870w, 1740s, 1680s, 1510s, 1455m, 1440m, 1390m, 1370w, 1350w, 1310m, 1280m, 1255m, 1150m, 1080w, 1050w, 1030w, 1000w, 945w, 910w, 700s. 1H-NMR (CDCl3): 7.37–7.22 (m, 10 arom. H, 1 NH); 6.19 (s, 1 NH); 5.45 (d, J = 6.9, 1 NH); 5.11, 5.06 (2d, JAB = 12.2, PhCH2O); 4.48–4.35 (m, HC(2) of Phe, HC(2) of Val); 3.68 (s, MeO); 3.16–3.02 (m, CH2 of Phe); 2.47–2.35, 2.25–2.12 (2m, 4 CH3, Me2CH of Val); 0.91, 0.88 (2d, J = 6.8, Me2CH of Val). 13C-NMR (CDCl3): 172.9, 172.3, 171.1 (3s, 3 C=O); 156.3 (s, OCON); 136.2, 135.8 (2s, 2 arom. C); 129.1, 128.9, 128.5, 128.3, 128.0, 127.3 (6d, 10 arom. CH); 67.3 (t, PhCH2O); 59.5 (s, C(4) of Tht); 57.3, 56.9 (2d, HC(2) of Phe, HC(2) of Val); 51.9 (q, MeO); 37.6 (t, CH2 of Phe); 33.7, 32.4 (2t, –CH2CCH2–); 30.9 (d, Me2CH of Val); 23.0, 22.9 (2t, –CH2CCH2–); 19.0, 17.7 (2q, Me2CH of Val). ESI-MS: 578 ([M+Na]+). [α]D –17.3 (c 1.013). Anal. Calcd for C39H49N3O7 (553.70): C 62.68, H 6.71, N 7.56, S 5.77. Found: C 62.68, H 6.80, N 7.56, S 5.70.

Z-Val-Tht-Phe-OEt (10d). The reaction of 497 mg (1.260 mmol) of 8e, 319 mg (1.389 mmol) 9b, 655 mg (1.260 mmol) PyBOP, and 520 μL (3.8 mmol) Et3N in 8 mL CH2Cl2 at rt, followed by CC (hexane/AcOEt 1:1) gave 637 mg (89%) of 10d. Colorless crystals; mp 157–158 °C. IR (CHCl3): 3660w, 3420m, 3060w, 3000m, 2960m, 2930m, 2870w, 1730s, 1680s, 1510s, 1455w, 1445w, 1390w, 1375w, 1350m, 1280m, 1220s, 1130m, 1110m, 1095m, 1080w, 1030w, 950w, 860w, 700s. 1H-NMR (CDCl3): 7.37–7.12 (m, 10 arom. H, 1 NH); 6.44 (s, 1 NH); 5.34 (d, J = 7.5, 1 NH); 5.10 (s, PhCH2O); 4.74 (q, J = 6.7, HC(2) of Phe); 4.08 (q, J = 7.1, MeCH2O); 3.91 (t, J = 7.3, HC(2) of Val); 3.14–3.00 (m, CH3 of Phe); 2.69–2.59, 2.50–2.39, 2.25–2.08 (3m, Me2CH of Val, 4 CH2); 1.17 (t, J = 7.1, MeCH2O); 0.96, 0.91 (2d, J = 6.8, Me2CH of Val). 13C-NMR (CDCl3): 172.7, 171.6, 171.4 (3s, 3 C=O); 156.7 (s, OCON); 136.1, 135.8 (2s, 2 arom. C); 129.3, 128.5, 128.3, 128.2, 128.0, 126.8 (6d, 10 arom. CH); 67.3 (t,
Reprotection of the piperidine N-atom of 8a.

Z-Phe-Pip(Z)-OH (11). To a solution of 75 mg (0.176 mmol) 8a in 2 mL 1M NaOH/dioxane at 0 °C were added 60 mg (0.352 mmol) benzyloxycarbonyl chloride (Z -Cl). The mixture was stirred at rt for 16 h, then 6M HCl was added until pH ≈ 2. The resulting mixture was extracted with AcOEt (2x), the organic phase was dried (MgSO4) and the solvent evaporated. Purification of the residue by CC (CH 2Cl2/MeOH 10:1) gave 37 mg (38%) of 11 as colorless resin. 1H-NMR ((D6)DMSO): 7.99 (s, 1 NH); 7.49 (d, J = 8.5, 1 NH); 7.39–7.15 (m, 15 arom. H); 5.07 (s, PhC \( \text{H}_2 \text{O} \)); 4.97, 4.92 (2d, J_{AB} = 12.8, PhC \( \text{H}_2 \text{O} \)); 4.40–4.25 (m, HC(2) of Phe); 3.70– 2.74 (m, –CH 2NCH2–, CH 2 of Phe); 2.15– 2.05, 1.80– 1.65 (2m, –CH 2CCH2–).

13C-NMR (CD3OD): 171.2 (s, 2 C=O); 156.1, 154.8 (2s, 2 OCON); 138.5, 137.5, 137.3 (3s, 3 arom. C); 129.6, 128.7, 128.6, 128.3, 128.1, 128.0, 127.8, 126.5 (8d, 15 arom. CH); 66.3, 65.6 (2t, 2 PhCH2O); 57.8 (s, C(4) of Pip); 56.8 (d, HC(2) of Phe); 40.4, 38.0 (2t, –CH2NCH2–, CH2 of Phe); 33.1, 32.2 (2t, –CH2CCH2–). CI-MS: 1185 (100, [2M–2H+3Na]+), 604 (27, [M–H+2Na]+).

General procedure for the selective deprotection of the carboxylic group of the tripeptide esters 10a–d. The corresponding tripeptide 10 was dissolved in THF/MeOH/H 2O (2:1:1) and cooled to 0 °C. Then, LiOH.H2O was added and the solution was stirred at rt for 2 h. After cooling to 0 °C, 2M HCl was added until pH ≈ 1 and the mixture was extracted with CH2Cl2 (3x). The combined organic phase was washed with 1M HCl, dried over MgSO4, and the solvent evaporated. Crystallization of the residue from CHCl3/hexane or CH2Cl2/Et2O gave the pure tripeptide acids 12.

Z-Phe-Thp-Val-OH (12a). The saponification of 150 mg (0.278 mmol) 10a in 4 mL THF/MeOH/H2O and crystallization from CH2Cl2/Et2O gave 133 mg (91%) of 12a. Colorless crystals; mp 132– 133 °C. IR (CHCl3): 3420 m, 3300 m, 3000 m, 2970m, 2930m, 2860m, 1715s, 1680s, 1515s, 1470m, 1455m, 1445m, 1390m, 1350m, 1300m, 1240m, 1150m, 1110m, 1080w, 1040w, 1030w, 890m, 700m. 1H-NMR (CDCl3): 7.38–7.18 (m, 10 arom. H); 6.62 (s, 1 NH); 5.80 (brx, 1 NH); 5.01 (s, PhCH2O); 4.55–4.48 (m, HC(2) of Phe); 4.41 (dd, J = 8.2, 4.9, HC(2) of Val); 3.69–3.57, 3.34–3.26, 3.14–3.06 (3m, –CH2OCH2–, CH2 of Phe); 2.23–2.04, 1.65–1.60 (2m, –CH2CCH2–, Me2CH of Val); 0.95, 0.88 (2d, J = 6.8, Me2CH of Val).

13C-NMR (CDCl3): 174.4, 173.4, 172.3 (3s, 3 C=O); 156.5 (s, OCON); 136.2, 135.8 (2s, 2 arom. C); 129.2, 128.9, 128.6, 128.4, 127.9, 127.3 (6d, 10 arom. CH); 67.3 (t, PhCH2O); 63.1, 63.0 (2t, –CH2OCH2–); 58.0 (s, C(4) of Thp); 57.8, 56.6 (2d, HC(2) of Phe, HC(2) of Val); 38.3, 33.2, 31.3 (3t,
CH$_2$ of Phe, –CH$_2$CCH$_2$–); 30.5 (d, Me$_2$CH of Val); 19.2, 17.7 (2q, Me$_2$CH of Val). ESI-MS: 548 ([M+1$^+$]). [α$^\text{D}$] –5.8 (c 1.010). Anal. Calcd for C$_{29}$H$_{35}$N$_3$O$_7$ (525.60): C 63.99, H 6.71, N 7.99. Found: C 63.68, H 6.70, N 7.99.

**Z-Val-Thp-Phe-OH (12b).** The saponification of 105 mg (0.190 mmol) 10b in 4 mL THF/MeOH/H$_2$O and crystallization from CHCl$_3$/hexane gave 91 mg (91%) of 12b. Colorless crystals; mp 155–157 °C. IR (CHCl$_3$): 3300m, 3030m, 2960m, 2880w, 1730s, 1670s, 1530m, 1500w, 1420m, 1390w, 1350m, 1280m, 1230m, 1170m, 1105m, 1030m, 830w, 780w, 740m, 700m. $^1$H-NMR (CD$_3$OD): 7.29–7.13 (m, 10 arom. H); 5.07, 5.03 (2d, $J_{AB}$ = 12.4, PhCH$_2$O); 4.58 (t, $J$ = 6.6, HC(2) of Phe); 3.92 (d, $J$ = 7.4, H(C2) of Val); 3.74–3.52, 3.52–3.34 (2m, –CH$_2$OCH$_2$–); 3.14–2.99 (m, CH$_2$ of Phe); 2.15–1.97, 1.88–1.84 (2m, Me$_2$CH of Val, –CH$_2$CCH$_2$–); 0.98, 0.96 (2d, $J$ = 6.6, Me$_2$CH of Val). $^{13}$C-NMR (CDCl$_3$): 174.9, 174.4, 174.3 (3s, 3 C=O); 158.6 (s, OCON); 138.1, 137.8 (2s, 2 arom. C); 130.3, 129.3, 129.2, 128.9, 128.7, 127.6 (6d, 10 arom. CH); 67.7 (t, PhCH$_2$O); 64.3, 64.1 (2t, –CH$_2$OCH$_2$–); 62.3 (d, HC(2) of Phe); 58.8 (s, C(4) of Thp); 55.2 (d, HC(2) of Val); 38.3 34.0, 31.9 (3t, CH$_2$ of Phe, –CH$_2$CCH$_2$–); 31.4 (d, Me$_2$CH of Val); 20.0, 18.9 (2q, Me$_2$CH of Val). ESI-MS: 1117 (46, [2M–2H+3Na]$^+$), 570 (32, [M–H+2Na]$^+$), 548 (100, [M+Na]$^+$). [α$^\text{D}$] –9.3 (c 1.009). Anal. Calcd for C$_{29}$H$_{35}$N$_3$O$_7$ (525.60): C 63.99, H 6.71, N 7.99. Found: C 63.91, H 6.91, N 8.03.

**Z-Phe-Tht-Val-OH (12c).** The saponification of 150 mg (0.270 mmol) 10c in 4 mL THF/MeOH/H$_2$O and crystallization from CHCl$_3$/hexane gave 145 mg (99%) of 12c. Colorless crystals; mp 102–103 °C. IR (KBr): 3390m, 3340m, 3140m, 3060m, 3015m, 2960m, 2930m, 1710s, 1695s, 1660s, 1650s, 1550s, 1540s, 1530s, 1505m, 1495m, 1450m, 1370w, 1360m, 1340w, 1310m, 1260m, 1240m, 1210m, 1155m, 1095w, 1070w, 1040m, 1030w, 1015w, 940w, 920w, 910w, 740m, 695m. $^1$H-NMR (CDCl$_3$): 7.95 (s, 1 NH); 7.44 (d, $J$ = 8.1, 1 NH); 7.35–7.20 (m, 10 arom. H, 1 NH); 5.08, 5.03 (2d, $J_{AB}$ = 12.5, PhCH$_2$O); 4.52–4.46, 4.26–4.20 (2m, HC(2) of Phe, HC(2) of Val); 3.04–2.95 (m, CH$_2$ of Phe); 2.63–2.44, 2.33–1.98 (2m, 4 CH$_2$, Me$_2$CH of Val); 0.92, 0.86 (2d, $J$ = 6.8, Me$_2$CH of Val). $^{13}$C-NMR (CD$_2$OD): 175.9, 174.7, 174.1 (3s, 3 C=O); 158.5 (s, OCON); 138.1, 138.0 (2s, 2 arom. C); 130.4, 129.7, 129.5, 129.0, 128.8, 128.1 (6d, 10 arom. CH); 67.7 (t, PhCH$_2$O); 61.1 (s, C(4) of Tht); 59.5, 58.0 (2d, HC(2) of Phe, HC(2) of Val); 51.9 (q, MeO); 38.6 (t, CH$_2$ of Phe); 35.8, 32.1 (2t, –CH$_2$CCH$_2$–); 31.7 (d, Me$_2$CH of Val); 23.9, 23.7 (2t, –CH$_2$SCH$_2$–); 19.6, 18.9 (2q, Me$_2$CH of Val). ESI-MS: 1106 (23, [2M+Na]$^+$), 564 (100, [M+Na]$^+$). [α$^\text{D}$] –13.1 (c 1.096). Anal. Calcd for C$_{29}$H$_{35}$N$_3$O$_7$S (559.24): C 60.09, H 6.66, N 7.56. Found: C 59.96, H 6.53, N 7.47.

**Z-Val-Tht-Phe-OH (12d).** The saponification of 171 mg (0.300 mmol) 10d in 4 mL THF/MeOH/H$_2$O and crystallization from CHCl$_3$/hexane gave 152 mg (94%) of 12d. Colorless crystals; mp 192–194 °C. IR (KBr): 3420s, 3020m, 2960m, 2940m, 2920m, 2870m, 1730s, 1710s, 1695s, 1670s, 1650s, 1555s, 1530s, 1515s, 1505s, 1475m, 1450m, 1430m, 1390m, 1370w, 1350m, 1330m, 1280s, 1210s, 1160m, 1140m,
1120 m, 1100 m, 1080 w, 1040 m, 980 w, 930 w, 855 m, 740 m, 695 m, 650 m. $^1$H-NMR (CD$_3$OD): 7.30–7.13 (m, 10 arom. H); 5.07 (s, PhCH$_2$O); 4.61–4.56, 3.96–3.92 (2m, HC(2) of Phe, HC(2) of Val); 3.15–2.98 (m, CH$_2$ of Phe); 2.84–2.71, 2.51–2.40, 2.26–1.97 (3m, Me$_2$CH of Val, 4 CH$_2$); 1.01, 0.96 (2d, $J = 6.8$, Me$_2$CH of Val). $^{13}$C-NMR (CDCl$_3$): 175.6, 174.5, 174.4 (3s, 3 C=O); 158.9 (s, OCON); 138.4, 138.0 (2s, 2 arom. C); 130.4, 129.5, 129.4, 129.0, 128.9, 127.7 (6d, 10 arom. CH); 67.8 (t, PhCH$_2$O); 62.7 (d, HC(2) of Phe); 60.9 (s, C(4) of Tht); 55.3 (d, HC(2) of Val); 38.4 (t, PhCH$_2$); 35.3, 33.2 (2t, –CH$_2$CCH$_2$–); 31.3 (d, Me$_2$CH of Val); 24.2, 23.9 (2t, –CH$_2$CH$_2$–); 20.1, 19.0 (2q, Me$_2$CH). ESI-MS: 1106 (100, [M+Na]$^+$), 564 (88, [M+Na]$^+$). [α$_D^+$] = 16.2 (c 1.077).

**General procedure for the selective deprotection of the amino group of the tripeptide esters 10a,b.**

To a solution of the corresponding tripeptide 10 in MeOH was added 0.1 mass-equiv. of Pd/C (10%). This mixture was stirred under an H$_2$-atmosphere at rt until disappearance of 10 (TLC). After filtration via Celite, the solvent of the filtrate was evaporated and the residue dried in HV.

*H-Phc-Thp-Val-OMe (13a).* Hydrogenolysis of 154 mg (0.285 mmol) of 10a in 5 mL MeOH at rt gave 115 mg (99%) of 13a. White crystals; mp 136–137 °C. IR (CHCl$_3$): 3420m, 3300m, 2960m, 1740s, 1675m, 1655s, 1540m, 1470m, 1455m, 1440m, 1390m, 1370m, 1300m, 1255m, 1190m, 1150m, 1110m, 1075w, 1050w, 1030m, 1010w, 990w, 970w, 930w, 700m. $^1$H-NMR (CDCl$_3$): 7.91 (d, $J = 8.3$, 1 NH); 7.79 (s, 1 NH); 7.37–7.23 (m, 5 arom. H); 4.47 (dd, $J = 8.3$, 5.1, HC(2) of Val); 3.83–3.75 (m, 2H of –CH$_2$OCH$_2$–); 3.72 (s, MeO); 3.64 (dd, $J = 9.8$, 3.8, HC(2) of Phe); 3.59–3.51 (m, 2 H of –CH$_2$OCH$_2$–); 3.32 (dd, $J = 13.8$, 3.8, 1 H of CH$_2$ of Phe); 2.72 (dd, $J = 13.8$, 9.8, 1 H of CH$_2$ of Phe); 2.34–2.16, 2.03–1.98 (2m, –CH$_2$CCH$_2$–, Me$_2$CH of Val); 0.96, 0.93 (2d, $J = 6.9$, Me$_2$CH of Val). $^{13}$C-NMR (CDCl$_3$): 175.5, 173.1, 172.3 (3s, 3 C=O); 137.5 (s, 1 arom. C); 129.3, 128.9, 127.1 (3d, 5 arom. CH); 66.44, 66.41 (2t, –CH$_2$OCH$_2$–); 57.6 (s, C(4) of Thp); 57.5, 56.8 (2d, HC(2) of Phe, HC(2) of Val); 52.0 (s, MeO); 40.7, 32.7, 32.5 (3t, CH$_2$ of Phe, –CH$_2$CCH$_2$–); 30.9 (d, Me$_2$CH of Val); 19.2, 17.8 (2q, Me$_2$CH of Val). CI-MS: 407 (20), 406 (100, [M+1]$^+$). [α$_D^+$] +17.3 (c 1.000). Anal. Calcd for C$_{21}$H$_{31}$N$_2$O$_4$ (405.49): C 62.20, H 7.71, N 10.36. Found: C 62.21, H 7.74, N 10.43.

*H-Val-Thp-Phe-OEt (13b).* Hydrogenolysis of 102 mg (0.184 mmol) of 10b in 2 mL MeOH at rt gave 72 mg (93%) of 13b. White foam. IR (CHCl$_3$): 3420w, 3300m, 3000m, 2970m, 2860m, 1740s, 1675s, 1510s, 1470m, 1455m, 1445m, 1390m, 1375m, 1350m, 1300m, 1240m, 1160m, 1110m, 1080m, 1030m, 860w, 820 w, 700m. $^1$H-NMR (CDCl$_3$): 7.87 (d, $J = 7.4$, 1 NH); 7.76 (s, 1 NH); 7.29–7.15 (m, 5 arom. H); 4.47 (q, $J = 7.5$, HC(2) of Phe); 4.14 (q, $J = 7.1$, MeCH$_2$O); 3.82–3.67, 3.62–3.48, 3.22–3.14, 3.05–2.97 (4m, –CH$_2$OCH$_2$–, HC(2) of Val, CH$_2$ of Phe); 2.36–2.21, 2.16–1.91 (2m, Me$_2$CH of Val, –CH$_2$CCH$_2$–); 1.22 (t, $J = 7.1$, MeCH$_2$O); 0.99, 0.81 (2d, $J = 6.9$, Me$_2$CH of Val). $^{13}$C-NMR (CDCl$_3$): 175.4, 172.8, 171.4 (3s, 3 C=O); 136.3 (s, 1 arom. C); 129.3, 128.2, 126.7 (3d, 5 arom. CH); 63.4, 63.3 (2t, –CH$_2$OCH$_2$–); 61.2 (t,
MeCH$_2$O); 61.1 ($d$, HC(2) of Phe); 57.2 ($s$, C(4) of Thp); 53.5 ($d$, HC(2) of Val); 37.8, 32.6, 32.2 ($3t$, CH$_2$ of Phe, –CH$_2$CH$_2$–); 30.3 ($d$, Me$_2$CH of Val); 19.7, 15.8 ($2q$, Me$_2$CH of Val); 14.0 ($q$, MeCH$_2$O).


X-Ray Crystal Structure Determination of 10a, 10c and 12a.$^{19}$ All measurements were made on a Rigaku AFC5R diffractometer using graphite-monochromated MoKα radiation ($\lambda = 0.71073$ Å) and a 12 kW rotating anode generator.$^{20}$ The intensities of three standard reflections were measured after every 150 reflections and remained stable throughout each data collection. The intensities were corrected for Lorentz and polarization effects.$^{21}$ Azimuthal scans of several reflections indicated no need for an absorption correction. Equivalent reflections, other than Friedel pairs, were merged. The data collection and refinement parameters are given below and views of the molecules are shown in Figures 2 and 4. Each structure was solved by direct methods using SHELXS-86,$^{22}$ which revealed the positions of all non-hydrogen atoms. The asymmetric unit of 12a contains one water molecule in addition to the peptide molecule. The non-hydrogen atoms were refined anisotropically. All of the amide, hydroxy and water H-atoms were placed in the positions indicated by difference electron density maps and their positions were allowed to refine together with individual isotropic displacement parameters. All remaining H-atoms were placed in geometrically calculated positions and refined using a riding model where each H-atom was assigned a fixed isotropic displacement parameter with a value equal to 1.2$U_{eq}$ of its parent C-atom (1.5$U_{eq}$ for the methyl groups). The refinement of each structure was carried out on $F^2$ by using full-matrix least-squares procedures, which minimized the function $\Sigma w(F_o^2-F_c^2)^2$. Corrections for secondary extinction were not applied. For 10c, refinement of the absolute structure parameter$^{23}$ yielded a value of –0.06(6), which confidently confirms that the refined model corresponds with the true enantiomorph. For the other structures, the absence of significant anomalous scattering elements meant that the precision of the absolute structure parameter was too low to be indicative; the enantiomer used in the refinement was based on the configuration expected from the chemical synthesis. Neutral atom scattering factors for non-H-atoms were taken from ref.$^{24}$, and the scattering factors for H-atoms were taken from ref.$^{25}$ Anomalous dispersion effects were included in $F_c$,$^{26}$ the values for $f'$ and $f''$ were those of ref.$^{27}$ The values of the mass attenuation coefficients are those of ref.$^{28}$ All calculations were performed using the SHELXL-2017$^{29}$ program.

Crystal data for 10a: C$_{29}$H$_{37}$N$_3$O$_7$, $M = 539.61$, crystallized from acetonitrile, colorless, prism, crystal dimensions 0.15 × 0.18 × 0.43 mm, monoclinic, space group $P2_1$, $Z = 2$, reflections for cell determination 25, 2θ range for cell determination 23–33°, $a = 6.137(3)$ Å, $b = 25.640(2)$ Å, $c = 9.561(2)$ Å, $\beta = 107.92(3)^\circ$, $V = 1431.4(9)$ Å$^3$, $T = 173(1)$ K, $D_x = 1.252$ g·cm$^{-3}$, $\mu$(MoKα) = 0.090 mm$^{-1}$, scan type $\omega$. 


$2\theta_{\text{max}} = 60^\circ$, total reflections measured 4633, symmetry independent reflections 4269, reflections with $I > 2\sigma(I)$ 3235, reflections used in refinement 4269, parameters refined 367, restraints 1, $R(F)$ [$I > 2\sigma(I)$ reflections] = 0.0453, $wR(F^2)$ [all data] = 0.1283 ($w = \left[\sigma^2(F_o^2)+(0.064P)^2+0.0064P\right]^{-1}$, where $P = (F_o^2+2F_c^2)/3$), goodness of fit 1.039, final $\Delta_{\text{max}}/\sigma$ 0.000, $\Delta\rho$ (max; min) = 0.25; –0.25 e Å$^{-3}$.

Crystal data for 10c: C$_{29}$H$_{37}$N$_3$O$_6$S, $M = 555.67$, crystallized from acetonitrile, colorless, prism, crystal dimensions $0.30 \times 0.30 \times 0.45$ mm, triclinic, space group $P1$, $Z = 1$, reflections for cell determination 25, $2\theta$ range for cell determination $39–40^\circ$, $a = 6.1404(10)$ Å, $b = 9.529(2)$ Å, $c = 13.951(3)$ Å, $\alpha = 71.994(15)^\circ$, $\beta = 89.050(18)^\circ$, $\gamma = 73.43(2)^\circ$, $V = 741.8(3)$ Å$^3$, $T = 273(1)$ K (lower temperatures damaged the crystals), $D_x = 1.244$ g·cm$^{-3}$, $\mu$(MoK$\alpha$) = 0.154 mm$^{-1}$, scan type $\omega/2\theta$, $2\theta_{\text{max}} = 55^\circ$, total reflections measured 7195, symmetry independent reflections 6784, reflections with $I > 2\sigma(I)$ 5398, reflections used in refinement 6784, parameters refined 367, restraints 3, $R(F)$ [$I > 2\sigma(I)$ reflections] = 0.0553, $wR(F^2)$ [all data] = 0.1634 ($w = \left[\sigma^2(F_o^2)+(0.1092P)^2+0.2482P\right]^{-1}$, where $P = (F_o^2+2F_c^2)/3$), goodness of fit 1.039, final $\Delta_{\text{max}}/\sigma$ 0.002, $\Delta\rho$ (max; min) = 0.39; –0.37 e Å$^{-3}$.

Crystal data for 12a: C$_{28}$H$_{35}$N$_3$O$_7$·H$_2$O, $M = 543.60$, crystallized from chloroform/hexane, colorless, plate, crystal dimensions $0.15 \times 0.42 \times 0.50$ mm, monoclinic, space group $P2_1$, $Z = 2$, reflections for cell determination 25, $2\theta$ range for cell determination $37–40^\circ$, $a = 6.159(4)$ Å, $b = 25.829(3)$ Å, $c = 9.502(3)$ Å, $\beta = 108.26(3)^\circ$, $V = 1435.6(10)$ Å$^3$, $T = 173(1)$ K, $D_x = 1.258$ g·cm$^{-3}$, $\mu$(MoK$\alpha$) = 0.092 mm$^{-1}$, scan type $\omega$, $2\theta_{\text{max}} = 55^\circ$, total reflections measured 3566, symmetry independent reflections 3363, reflections with $I > 2\sigma(I)$ 3070, reflections used in refinement 3363, parameters refined 378, restraints 1, $R(F)$ [$I > 2\sigma(I)$ reflections] = 0.0336, $wR(F^2)$ [all data] = 0.0845 ($w = \left[\sigma^2(F_o^2)+(0.0392P)^2+0.2482P\right]^{-1}$, where $P = (F_o^2+2F_c^2)/3$), goodness of fit 1.017, final $\Delta_{\text{max}}/\sigma$ 0.000, $\Delta\rho$ (max; min) = 0.18; –0.18 e Å$^{-3}$.

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REFERENCES AND NOTES

1. Part of the PhD Thesis of C. S., University of Zurich.


CCDC-1815720–1815722 contain the supplementary crystallographic data for this paper. The data can be obtained free of charge from the Cambridge Crystallographic Data Centre via www.ccdc.cam.ac.uk/structures.


