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Asymmetric Synthesis of New \(\beta\)-Lactam Lipopeptides as Bacterial Signal Peptidase I Inhibitors

Céline Crauste,[a] Matheus Froeyen,[a] Jozef Anné,[b] and Piet Herdewijn*[a]

Keywords: Medicinal chemistry / Antibiotics / Enzymes / Peptides / Lactams / Asymmetric synthesis

The transmembrane bacterial enzyme, signal peptidase I, is recognized as being a promising target for reducing the emergence of drug resistance. The asymmetric synthesis and the biological evaluation of original \(\beta\)-lactam lipopeptides have been performed to discover potent signal peptidase inhibitors. The importance of the azetidinone motif of these lipopeptides has been demonstrated and can serve as a starting point to exploit and improve the reactivity of the \(\beta\)-lactam in peptidomimetics.

Introduction

Faced with the emergence of drug-resistant bacteria, the discovery of antibiotics that act by novel and various mechanisms is of crucial importance. The bacterial enzyme, signal peptidase I (SPase I), has received increasing interest as a potential new target for the discovery of antibacterial agents.[1] This transmembrane enzyme appears to be essential for the viability of both Gram-positive and -negative bacteria (i.e., *Escherichia coli* and *Staphylococcus aureus*).[2,3] Its physiological function is to ensure the release of secreted bacterial proteins by cleaving off the signal peptide from translocated pre-proteins. The bacterial signal peptidases use a unique dyad mechanism that involves serine and histidine amino acids.[4] As a consequence, it should be possible to develop specific SPase I inhibitors with high selectivity and low toxicity.

The natural substrates of SPase I contain a C-terminal domain (c-region) that is responsible for substrate recognition. In most cases alanines are found in proximity to the cleavage site in the P1 and P3 positions.[5,6] Therefore the substrate recognition rule is often called the Ala-X-Ala (AXA) rule. This observation was confirmed by the crystal structure of *E. coli* SPase I LepB, with two small hydrophobic pockets (S1 and S3) identified on the surface of the catalytic domain of the enzyme in which both alanine residues (AXA) of natural peptide substrates are oriented.[7]

Various classes of molecules have been described as potential inhibitors of SPase I of *E. coli*: \(\beta\)-Lactam compounds such as the allyl (5S,6S)-6-[(R)-acetoxyethyl]penem-3-carboxylate (Penem A, see part A of Figure 1 and Figure 2),[8,9] peptides (Arylomycin A2)[10] and lipoglycopeptide derivatives.[11] For peptide derivatives, N-terminal acylation with fatty acids like decanoyl acid seem to enhance the potency of the inhibitors.[12,13] The decanoyl chain is believed to interact with the membrane or the detergent used for modeling the membrane bilayer in in vitro assays.[7]

Up to now, small peptides that mimic the protein cleavage site have proven to be inefficient as SPase I inhibitors.[14] However, we first investigated the possibility of developing small peptidomimetics as potential SPase I inhibitors based on the presence of the \(N\)-decanoyl side-chain and the AXA moiety. Our work[13] led to the discovery of a decanoyl-PTANA-COH peptide aldehyde (Figure 2) with an IC\(_{50}\) value of 0.09 \(\mu\)M when tested against SPsB, the SPase I of *S. aureus*. This lipopeptide inhibitor has a C-terminal aldehyde function that allows nucleophilic attack of the serine amino acid involved in the peptide cleavage mechanism.[15,16] Despite its “serine trap” efficiency, the aldehyde function is often metabolically unstable and hence its replacement by other classes of electrophiles would be a promising optimization. Several chemically reactive groups have been introduced into protease inhibitor structures to act as “serine traps”, like trifluoromethyl ketones,[17] \(\alpha\)-keto amides,[18] \(\beta\)-lactams,[19,20] or boronic acids.[21,22] Among them, the \(\beta\)-lactam residue is the only one previously described in non-peptidic SPase I inhibitors.[8,9] Therefore, to improve the activity of the decanoyl-PTANA-COH lipopeptide and to obtain more information on the binding of this kind of inhibitor in the active site of the enzyme, we replaced the aldehyde function by \(\beta\)-lactam residues, commonly used in other protease inhibitors.[20,23,24] (compounds...
Figure 1. A: Structure of Penem A in *E. coli* SPase I covalently bound to Ser90.[7] B: Model of decanoyl-PTAN-aminoethyl-azetidinone 1a (without decanoyl) in the *E. coli* SPase I active site. The S1 site is printed in yellow, the S3 site in red, and common residues in orange. The Ser90 and Lys145 active site residues are also shown (C atoms: grey). The two alanine side-chains in the AXA motif are printed in cyan.

Described Spase I inhibitors

Decanolic peptide aldehyde: Decanolic-PTAN-COH

New β-lactam lipopeptide Spase I inhibitors

1a: (3S,5R)
1b: (3S,5S)
2a: (5S,6R)
2b: (5S,6S)
2c: (3S,5S)

Synthetic precursors

Figure 2. Chemical structures of previously described SPase I inhibitors (decanoic-peptide-aldehyde[13] and Penem A[8,9]), potential new lipopeptide inhibitors (1 and 2), and their synthetic precursors.

1 and 2). Because an appropriate stereochemistry of the β-lactam structure is essential for its biological activity,[8,25]

we focused our work on the synthesis of several isomers of this new class of β-lactam lipopeptides. We report herein the design, synthesis, and biological evaluation of these new β-lactam-lipopeptides.

Results and Discussion

The research started with modeling experiments by docking potential peptidomimetics in the catalytic site of the enzyme and selecting the best-fitting molecules for future synthesis. The model of the new lipopeptides was created starting from the X-ray structure of the *E. coli* Signal peptidase I acyl-enzyme[7] covalently bound to the 5,6β-lactam inhibitor Penem A (Figure 2). This crystal structure (Figure 1, A) shows that the binding to *E. coli* SPase I can be influenced by the properties of the C6 substituent of such penems oriented towards the S1 hydrophobic pocket, whereas modification at the C2 and C3 positions offers little or no opportunity for binding optimization.[26] In fact, co-crystallization reveals that the 16-methyl group of Penem A is oriented toward the S1 pocket of the active site of the enzyme (like the alanine residue in the P1 position of natural substrates). However, no access to the S3 pocket is revealed with this penem inhibitor.

As the modeling experiments[7] predicted that the methyl moiety of a β-lactam alkyl ester residue could mimic the last amino acid of the AXA sequence oriented towards the S1 pocket, we investigated the design of a new class of peptide inhibitors to optimize the orientation towards both the S1 and S3 pockets. We planned to combine the structures and properties of the decanoyl-peptide and a β-lactam residue. Therefore we decided to replace the last alanine residue of the decanoyl-PTANA-COH peptide by the aminoethyl-azetidinone 9 (Figure 2) to create lipopeptide of type 1 with a monocyclic β-lactam moiety. We initially chose a monocyclic β-lactam (mono-bactam) residue because no interactions were observed between the enzyme and the five-membered ring of Penem A.[26] By introducing the decanoyl-PTAN moiety onto the amino group of the azetidinone 9, we expected to optimize the interactions with the S3 pocket to increase the affinity for the active site.
β-Lactam Lipopeptides as Bacterial Signal Peptidase I Inhibitors

To choose the appropriate stereochemistry for the synthesis of the lipopeptide 1, we first focused on the results published by Allsop et al. [8] According to them, the 15R configuration of Penem A seems to be essential for activity. Indeed by modeling both lipopeptides 1a and 1b inside the E. coli SPase I active site, a better fit was observed for the 5R lipopeptide 1a (Figure 1, B). [27] In this theoretical model, the two methyl functions (the methyl group of alanine and the 6-methyl group) colored in cyan appear to fit into both hydrophobic pockets S1 and S3. However, the C5 carbon of the lipopeptide 1a, which mimics the α carbon of the alanine residue, has an R configuration (o), in contrast to the natural S configuration (t) seen for all natural amino acids. Because of this observation both the 5R and 5S isomers of molecule 1 were synthesized and evaluated.

Based on the structures of the two SPase I inhibitors described (Figure 2), we also explored the direct coupling of the decanoic peptide with the commercially available and inexpensive 6-aminopenicillanic acid (6-APA, compounds 2a-d). In this case, substrate recognition was based only on the decanoyl-PTAN moiety and the “serine trap” reactivity was introduced through a bicyclic penam moiety. The biological activity of the Penem A has been described only for its cis-SS,6S isomer, which has the opposite stereochemistry to most β-lactam antibiotics, and the starting material 6-APA. Thus, synthetic schemes had to be elaborated to invert both stereocenters of the 6-APA to obtain the four diastereoisomers of compound 2 and explore the importance of the stereochemistry for the activity of this new class of potential SPase I inhibitors. Indeed, synthetic schemes to invert the configuration and obtain the amine-free penam of the 6-APA derivative are poorly described due to the instability of the unprotected penam β-lactam ring.

The synthesis of the lipopeptides 1 and 2 first required the solid-phase synthesis of decanoyl-PTAN-COOH peptide 4, which was then coupled to the appropriate residues 9 or 14. The supported synthesis was performed on a Wang resin using classical coupling reagents (HOBt/DIPEA/DIC) for peptide bond formation and Fmoc-protected amino acids. [29] The last steps involved N-terminal decanoylation followed by side-chain deprotection and cleavage from the solid support using a TFA/H₂O/thioanisole solution.

The azetidinone-peptides 1a, b were synthesized by condensation between decanoyl-PTAN-COOH and the (5R)- or (5S)-(1-aminoethyl)azetidinone 9a or 9b (for atom numbering see Scheme 1). We used the commercially available silylated azetidinone as the starting material to give the (5R)-hydroxyazetidinone 6a (Scheme 1). Reduction of the commercially protected β-lactam with NaBH₄ in ethanol at 10 °C [30] led to the C4-unsubstituted compound 5 in 81% yield. Deprotection of the silyl group was performed by using HCl solution [30] to afford 6a in 77% yield whereas TBAF deprotection [31] gave only 40% yield due to some degradation. The hydroxyazetidinone 6a obtained was then used as the starting material to synthesize both the (5R)- and (5S)-aminoazetidinones 9a and 9b.

The (5S)-(1-aminoethyl)azetidinone 9a was synthesized following a sequence of mesylation, aziridination, and reaction processes. Unexpectedly, the trivial conversion of the hydroxy function into an activated mesylate 7a presented some difficulties. [32] Of the solvents tested (DMF, DCM, and pyridine), only pyridine was found to be appropriate and several additions of MsCl during the reaction were needed to achieve completion of the reaction. Nucleophilic displacement of the mesylate 7a with lithium azide was performed at 60 °C to provide the (5S)-azidoazetidinone 8a (81% yield), which, after hydrogenolysis, afforded the desired aminoazetidinone 9a.

To obtain the 5R isomer 9b from the hydroxazetidinone 6a, an inversion of the absolute configuration at the C5 position was performed by Mitsunobu esterification followed by saponification. [33,34] Under the usual Mitsunobu conditions of DEAD (diethyl azodicarboxylate), PPh₃ (triphenylphosphane), and benzoic acid, the reaction was low yielding and most of the starting material was recovered. Several reaction methods were investigated in an attempt to improve its efficiency. [53] As this reaction is known to be highly sensitive to the steric hindrance of the starting alcohol, several methods were developed to overcome this drawback. We tried to perform the nucleophilic substitution by replacing benzoic acid by a stronger acid (nitrobenzoic acid). However, additional degradation reactions and the formation of side-products were observed. The activation
of the hydroxy function was then tested by using a mixture of DIAD (diisopropyl azodicarboxylate)/PPh₃, ADDP (azodicarboxyldipiperidine)/PBu₃ (tributylphosphane) or TMAD (N,N,N',N'-tetramethylazodicarboxamide)/PBu₃. The use of ADDP or TMAD as activating reagent increased the yield up to 35%. These conditions also facilitated the purification step. (5S)-Hydroxyazetidinone 6b was then obtained after saponification. Finally, the desired (5R)-aminoazetidinone 9b was obtained from the hydroxy derivative 6b after mesylation, azidination, and reduction processes, as described above.

Both (5R)- and (5S)-aminoazetidinones 9 were then coupled to the previously synthesized decanoyl-PTAN-COOH (4) by using the usual coupling reagents (HOBt/EDCI). The 5S isomer of the lipopeptide 1 was obtained in diasteromerically pure form. However, after purification by chromatography and semi-preparative reversed-phase HPLC, 1H NMR analysis of the 5R isomer 1a showed a small proportion of a second product, probably formed by epimerization at the C3 position.

In the second part of our work, we focused on the synthesis of lipopeptides 2a-d, directly conjugated to penam residues. This synthetic pathway started with the synthesis of the four 5,6 diastereoisomers of the commercially available 6-APA (14a-d; Scheme 2). Epimerization at the 6- and/or 5-positions has already been reported for penicillin bearing an imido substituent at the 6-position and ester protection of the carboxylic function. Indeed, it has been recognized that phthalimido-substituted β-lactam is more stable to a wide variety of reaction conditions than the corresponding amido derivatives. Moreover, the synthetic yield of the cis-5S,6S isomer 14d, starting from the (5R,6R)-6-APA, seems to be improved when a phthalimido-protected strategy is used.

To perform the epimerization steps, the sodium salt of commercial 6-APA was treated with N-ethoxycarbonylphthalimide and then esterified in the presence of benzyl bromide and triethylamine to obtain the protected cis-5R,6R isomer 11a in 42% yield (for the two steps; Scheme 2). From compound 11a, epimerization at the 6-position was accomplished by using a catalytic amount of the strong base 1,8-diazabicyclo[5.4.0]undec-7-ene (DBU) to give the trans-5R,6S isomer 11b in good yield (80%). Then the Kukolja protocol was employed to perform the epimerization at the 5-position. From the trans isomer 11b, 1 equiv. of sulfuryl chloride was used to cleave the C5–S bond of the penicillin thiazolidine ring. The trans and cis chloro derivatives were obtained in a ratio of 9:1. The mixture was then treated with anhydrous tin(II) chloride to achieve the highly trans-selective recyclization of the second cycle of the penam system. Thus, a mixture of both cis-(5S,6S)-11d (31% in two steps) and trans-(5R,6S)-11b isomers were obtained and separated by silica gel chromatography. Application of the two-step Kukolja protocol to the cis-5R,6R isomer 11a yielded the trans-5S,6R isomer 11c without any recovery of the starting isomer.

A last deprotection step had to be performed to realize the coupling to the decanoic peptide. Several examples of successful dephthaloylation in one step using hydrazine derivatives have been reported for monocyclic β-lactam, but only a few examples are described for bicyclic derivatives like cephalosporins, caracephems, and isocaracephems. As previously reported, we observed that it was not possible to carry out efficient hydrazinolysis on penicillin derivatives without β-lactam ring-opening. The reaction performed at a lower temperature (−25 to −78 °C) or by using a less reactive hydrazine (N-methylhydrazine) preserved the β-lactam ring but decreased the efficiency of the deprotection. To overcome this instability of the β-lactam, the phthalimido group had to be converted into a more unstable phthalisothioimido group prior to hydrazinolysis. For the first time this “three-step” method was applied separately to the deprotection of the four diastereoisomers 11a-d (Scheme 3).

Compounds 11a-c were hydrolyzed to give the phthalimide acids by slow addition of 1 equiv. of aqueous sodium sulfide at 0–5 °C over 15 min (48–67%; Scheme 3). Under the same basic conditions, cis-(5S,6S)-11d appeared to be much more sensitive and unstable compared with the other isomers, probably because of a bigger strain in the bicyclic ring. To limit the degradation of the β-lactam ring, less than 1 equiv. of sodium sulfide was used as well as a careful neutralization. The phthalimide acids 12 were cyclized to phthalisothioimides 13 by using N,N-dicyclohexylcarbodiimide as the activating reagent. Only the phthalisothioimide form (13a-c) was observed by 1H NMR analysis in both the crude materials and the purified compounds (45–70%) without any traces of the phthalimide form. However, during the synthesis of the cis-(5S,6S)-13d, purification by silica gel chromatography led to the total conversion to the starting...
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The inhibitory activities of the synthesized compounds were evaluated by a continuous fluorimetric assay.[13,46] This quantitative in vitro assay is based on the processing of an internally quenched fluorescent synthetic peptide substrate (FRET-based assay). After cleavage of the peptide by the SPase, an increase in the fluorescence was measured. Therefore the inhibition of SPase I can be analyzed and quantified in the presence and absence of the synthetic lipopeptides. The results of the in vitro assays on the isolated enzyme are summarized in Table 1.

Table 1. Results of the inhibitory activity assay of the synthesized lipopeptides on S. aureus.[a]

<table>
<thead>
<tr>
<th>Entry</th>
<th>Compound</th>
<th>Conc. [µM]</th>
<th>Inhibition [%]</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>B[10]</td>
<td>200</td>
<td>10</td>
</tr>
<tr>
<td>2</td>
<td>1a</td>
<td>100</td>
<td>57</td>
</tr>
<tr>
<td>3</td>
<td>1b</td>
<td>100</td>
<td>47</td>
</tr>
<tr>
<td>4</td>
<td>2a</td>
<td>100</td>
<td>25</td>
</tr>
<tr>
<td>5</td>
<td>2'a (COOH)[5]</td>
<td>100</td>
<td>21</td>
</tr>
<tr>
<td>6</td>
<td>2b</td>
<td>100</td>
<td>36</td>
</tr>
<tr>
<td>7</td>
<td>2c</td>
<td>100</td>
<td>36</td>
</tr>
<tr>
<td>8</td>
<td>2d</td>
<td>100</td>
<td>26</td>
</tr>
<tr>
<td>9</td>
<td>3 (decanoyl-PTAN-NH₂)</td>
<td>100 × 10⁻³</td>
<td>60</td>
</tr>
</tbody>
</table>

[a] Concentration of the SPsB Spase I: 0.25 µM; concentration of the quenched peptide: 5 µM. Experiments were performed in duplicate. [b] Ref.47 [c] Deprotected form of 2a.

The biological evaluation showed the lipopeptides 1 and 2 to be moderately-to-weakly active. In each case we found that the modifications introduced into the lipopeptide did not reach the inhibition observed for the lipopeptide aldehyde (decanoyl-PTANA-COH). However, the high activity observed for the new lipopeptides 1 and 2 compared with the previously studied hydroxyethyl isoster B[47] (entry 1, Table 1 and Figure 3) confirms that the introduction of a β-lactam moiety at the carboxylate end seems to be a better approach to obtaining potent SPase I inhibitors. The results obtained for the β-lactam inhibitors show slight differences depending upon the compound. Indeed, the percentage inhibition obtained for the mono-bactam compounds (entries 4–8) appears to be two-fold higher than those obtained for the penicillin peptides (entries 4–8; 25, 21, 36, 12, and 26%). This result does not correspond to the chemical reactivity of either the mono-bactam or penam as the amide function of the bicyclic derivative is usually more reactive. However, we did not consider the time frame of the inhibition experiments during the first screening assay and this study may be the subject of further investigation.

![Figure 3. Additional compounds tested in the FRET-based assay on S. aureus sPsB.](image-url)
To eliminate the influence of solubility problems on the small differences of inhibitory activity, the deprotected carboxylic acid of 2a was synthesized (by hydrogenation) and evaluated (2′a, entry 5). The same percentage inhibition was obtained for both the protected and the deprotected compounds. The methyl group used as a mimic of the last alanine of the AXA sequence (present only in compounds 1a and 1b) might play a role in the recognition of the active site. To demonstrate the importance of the β-lactam moiety in the inhibition observed, decanoyl-PTAN-NH₂ (3; Figure 3) was also synthesized from peptide 4 (by treatment with ammonium chloride) and tested on SPase I SpsB. To reach the same level of inhibition as compounds 1, decanoyl-PTAN-NH₂ (3; devoid of the β-lactam moiety) had to be tested at 100 μm, a concentration 10³-fold higher than that used for the lipopeptides 1. As the removal of the β-lactam moiety resulted in a complete loss of activity, this function might be implicated in the binding to the active site. However, regarding the influence of the stereochemistry of compounds 1 and 2, no significant differences were observed between the activities of the several isomers synthesized. As a consequence, the penam or the mono-bactam residue might not be ideally accommodated in the active site and for this reason only modest activities were observed with compounds 1 and 2.

Inhibition tests were also performed on the SPase I enzyme best characterized until now, the LepB of E. coli, the crystal structure of which was used in the modeling experiment (as the S. aureus SPase I has not yet been crystallized). No significant activities were observed at 10 μm. Indeed the decanoyl-PTANA sequences appear less suitable for binding to E. coli SPase I than to the enzyme of Gram-positive species. However, a structural study should be performed to find the appropriate peptide sequences that would allow both to adopt a β-sheet-like conformation and to fit correctly the β-lactam ring into the active site of the SpsB enzyme. Furthermore, given the large differences in the biochemical activity of the peptide aldehyde[13] and the present β-lactam conjugates, it might be hypothesized that, in addition to the non-optimal positioning in the active site, the reactivities of the β-lactam moieties of compounds of type 1 and 2 are not high enough to allow covalent bond formation. A β-lactam lipopeptide obtained from coupling between the decanoic-peptide and a penam moiety like Penem A could be a better alternative to achieve higher reactivity.

Conclusions

A new class of lipopeptides bearing a β-lactam moiety has been synthesized. This class of compounds has been poorly described and usually only the racemic form of the penicillin moiety is represented.[48] An effort has been made to use stereoselective procedures and to obtain several diastereoisomers of the lipopeptides 1 and 2. The deprotection of the phthalimide group was performed for the first time on the four diastereoisomers of the protected 6-APA residue, opening the way for further asymmetric synthesis of penam compounds. In vitro studies showed moderate SPase I inhibition, which could be due to inappropriate positioning of the β-lactam in the active site. However, because the β-lactam motif seems to be involved in the inhibitory activity, other peptide sequences bearing the PTANA motif should be envisaged, as well as modifications of the β-lactam moiety, to improve the binding and reactivity with the enzyme SpsB and for the design of new potential peptidomimetic β-lactam lipopeptides.

Experimental Section

General: NMR spectra were recorded with a Bruker UltraShield 300 (1H: 300 MHz; 13C: 75 MHz; 31P 121 MHz), 500 MHz (1H: 500 MHz) or 600 MHz (1H: 600 MHz) spectrometer using tetramethylsilane as internal standard. Chemical shifts are reported in δ units and coupling constants (J values) in Hz. Reagents and solvents were obtained from commercial sources (Sigma–Aldrich, Acros-Organics, NovaBiochem, Alfa-Aesar and TCI Europe) and used as received (THF and DCM were used after distillation over Na/benzophenone and CaH₂, respectively). The progress of the reaction was monitored either by TLC, performed on Alugram SIL G/UV₂₅₄ plates, or by mass analysis performed with a TSQ Quantum Thermo Scientific instrument. Column chromatography was performed with Grace Silicagel DAVISIL LCA (0.040–0.063 or 0.070–0.200 mm). Purification on HPLC was realized with a semi-preparative Polymer Lab column, PRLPS 10 μm (21.2 cm × 150 mm ID; mobile phase flow; 2 mL/min, gradient H₂O/CH₃CN or H₂O/CH₃CN/0.1% TFA, UV detection 214/254 nm). Optical rotations were measured with a Perkin–Elmer polarimeter (model 341) in chloroform. All final lipopeptide purities were checked by HRMS and HPLC. HRMS spectra were recorded with a Q-ToF-2 (Micromass, Manchester, UK) mass spectrometer. Accurate masses were determined by co-infusion of the samples with 2 μm leucine enkephalin (YGGFL) and recalibration of the spectrum using the peak at m/z = 556.2771 as the lock mass. At least 10 spectra were recorded and averaged. HPLC was performed with a Shimadzu SPD-M20A HPLC System using an intersil ODS-3 4 μm (4.6 cm × 100 mm ID) column. The compounds were analyzed with H₂O/CH₃CN/0.1% TFA or H₂O/CH₃CN solvent systems with a 30 or 35 min gradient: 5–100% CH₃CN with a flow rate of 1.0 mL/min.

Abbreviations: HOBt: hydroxybenzotriazole; DIPEA: N,N-diisopropylamine; DIC: N,N-diisopropylcarbodiimide; DMF: dimethylformamide; TFA: trifluoroacetic acid; EDCl: 1-ethyl-3-(3-dimethylaminopropyl)carbodiimide; DCM: dichloromethane; 6-APA: 6-aminopenicillanic acid; PTAN: 1-prolyl-1-threonyl1-ala-nyl-1-asparaginyl.

Peptide Synthesis

Decanoyl-1-prolyl-1-threonyl1-ala-nyl-1-asparaginyl (4): The solid-phase synthesis of this peptide was performed on Wang resin (0.80 mg, 0.52 mmol) following a Fmoc strategy. DMF was used as solvent, HOBt (4 equiv.), DIPEA (2 equiv.), and DIC (4 equiv.) as coupling reagents, and all the amino acids coupled were Fmoc-protected. First, Fmoc-asparagine(trt) (4 equiv.) was coupled to the resin over 5 h. After capping in the presence of acetic anhydride (20% in pyridine) and Fmoc deprotection in the presence of piperidine (20% in DMF), the next amino acid was coupled. The same process was performed for the coupling of Fmoc-alanine (4 equiv.), Fmoc-threonyl(tbu) (4 equiv.), Fmoc-proline (4 equiv.), and then...
decanoic acid (4 equiv.). The protec-tion of the lateral protecting groups and the cleavage of the resin were performed with a solution of TFA/thiophene/H2O (9.5:0.25:0.25) over 3 h. The desired peptide 4 (62 mg, 21% in steps) was then obtained after precipitation in diisopropyl ether and purification by preparative reversed-phase HPLC (A: H2O/0.1% TFA/B: CH3CN/0.1% TFA). 1H NMR (300 MHz, CDCl3): δH = 4.48 [m, 1 H, H2O], 2.43–2.10 [m, 4 H, H2O, Tr, H(N)] ppm. 13C NMR (75 MHz, CDCl3): δC = 167.5, 165.6, 132.6, 129.7, 129.3, 131.7, 173.3, 173.1, 172.9, 172.5, 171.2, 66.3, 60.1, 58.1, 49.7, 49.5, 35.9, 34.0, 31.2, 28.8, 28.7, 28.6, 24.3, 24.1, 21.9, 18.6, 15.7, 13.2 ppm. HRMS (EI): calcld. for C24H42NO8 [M + H]+ 556.3346; found 556.3354.

Decanoyl-l-prolyl-l-threonyl-l-alanyl-l-asparaginylcarboxamide (3): A solution of the acid (10.0:10.0:0.5) in ethyl acetate was added to a solution of the azetidinone 6a (50 mg, 0.14 mmol) in dry pyridine at 0 °C. After filtration, the reaction mixture was washed with ethyl acetate and several times with water and brine. The aqueous phase was extracted several times with DCM. The organic phases were combined and the solvents were removed under reduced pressure. The residue obtained was purified by chromatography on silica gel using ethyl acetate/MeOH (95:5) as eluent to give 6b (35 mg, 35%) as a white amorphous solid. HRMS: calcd. for C21H31NO4 [M + H]+ 327.2396; found 327.2405.

Synthesis of Peptide Type 1

(3S)-(3R)-(1R)-(tert-Butyldimethylsilyloxy)ethylazetidin-2-one (5): NaBH4 (0.72 g, 16.10 mmol) was added to a solution of the commercial silylated azetidine (5 g, 17.40 mmol) in absolute ethanol (25 mL) at 0 °C to cool the reaction solution. The suspension was kept at 10 °C for 1 h 30 min. After filtration, the reaction mixture was washed with ethyl acetate and several times with DCM. The filtrates were combined and the solvents were removed under reduced pressure. The residue obtained was purified by chromatography on silica gel using ethyl acetate/MeOH (95:5:5) as eluent to give 5 (5.00 g, 71%) as a white solid. HRMS: calcd. for C22H35NO3 [M + H]+ 367.2664; found 367.2672.

(3S)-[3R]-[1R]-Hydroxyethylazetidin-2-one (6a): Compound 5 (0.50 g, 2.20 mmol) was dissolved in CH2CN (12 mL), and 1 N HCl (2 mL) was added. The reaction was monitored by TLC and after 1 h 30 min another 1 N HCl (0.50 mL) was added to avoid total deprotection. After 30 min, the mixture was neutralized by the slow addition of a 1 N NaOH solution. The solvents were removed under reduced pressure and the residue obtained was purified by chromatography on silica gel using DCM/ethyl acetate/MeOH (10:10:0.3) as eluent to give 6a (194 mg, 77%) as a white solid. HRMS: calcd. for C12H20NO3 [M + H]+ 212.1370; found 212.1371. 1H NMR (300 MHz, CDCl3): δH = 4.04 (br., 1 H, NH), 3.75 (br., 1 H, NH), 3.43 (br., 1 H, NH), 3.15 (br., 1 H, NH), 3.06–3.04 (m, 2 H, 4-H, 3-H) ppm. 13C NMR (75 MHz, CDCl3): δC = 167.9, 64.5, 58.2, 37.6, 22.1 ppm. HRMS (EI): calcld. for C12H20NO3 [M + H]+ 212.1371; found 212.1371.

(3S)-[3S]-[1S]-[1R]-Hydroxyethylazetidin-2-one (6b): Tributylphosphine (0.16 mL, 0.65 mmol) and benzoic acid (53 mg, 0.43 mmol) were added at 0 °C to a solution of the azetidinone 6a (50 mg, 0.14 mmol) in dry toluene (3 mL). Then, N,N,N′,N′-tetramethyl-azidocarboxamide (TMAD; 112 mg, 0.65 mmol) was added in one portion. The reaction mixture was stirred at 0 °C for 10 min and then at 60 °C for 1 h. Dihydro-TMAD was precipitated and filtered off after the addition of hexane to the reaction mixture. The solvents were removed under reduced pressure and the resulting residue was purified by chromatography on silica gel using hexane/ethyl acetate (8:2:1) as eluent to give 6b (35 mg, 35%) as a white amorphous solid. HRMS: calcd. for C21H31NO4 [M + H]+ 327.2396; found 327.2405.
(3S)-3-[(1S)-1-(Methylsulfonyloxy)ethyl]azetidin-2-one (7b): The same process was used starting from the alcohol 6b (30 mg, 0.26 mmol) and afforded the mesylate 7b (30 mg, 60%) as a white solid after purification by chromatography on silica gel using hexane/ethyl acetate (2:8) as eluent. Rf = 0.3 (ethyl acetate). 1H NMR (300 MHz, CDCl3): δH = 5.98 (br., 1 H, NH), 5.12 (quint., J = 6.5 Hz, 1 H, 5-H), 3.57 (m, 1 H, 3-H, 4-H), 3.45 (t, J = 6.0 Hz, 1 H, 4-Hb), 3.35 (dd, J = 2.5, J = 6.0 Hz, 4-Ha), 3.10 (s, 3 H, CH2SO2), 1.62 (d, J = 6.5 Hz, 3 H, 6-H) ppm. 13C NMR (75 MHz, CDCl3): δC = 166.3, 74.9, 55.3, 38.6, 37.7, 18.4 ppm. HRMS (EI): calcd. for C8H14NO4S [M + H]+ 194.0486; found 194.0494.

Synthesis of Lipopeptides Type 2

(3S)-3-[(1R)-1-Azaglycyl]azetidin-2-one (8a): LiN3 (8.07 mg, 0.16 mmol) was added to a solution of the mesylate 7a (22 mg, 0.06 mmol) in dry DMF at room temperature. The reaction mixture was stirred overnight at 60 °C and the solvent was evaporated under reduced pressure. The residue obtained was dissolved in DCM and washed with water and brine. The aqueous phase was combined, dried (MgSO4), and concentrated under vacuum. The residue obtained was purified by chromatography on silica gel using hexane/ethyl acetate (9.5:0.5 to 8:2) as eluent to give 1a (11 mg, 62%) as a white solid contaminated by the 3R isomer. Major product: Rf = 0.5 (DCM/MeOH, 9:1). 1H NMR (300 MHz, CDCl3): δH = 4.62 (m, 1 H, Hα), 4.46–4.20 (m, 4 H, Hβ(Ala, Pro, Thr), Hβ(Thr)), 3.69, 3.54 (m, 2 H, CH2β(Pro)), 3.45–3.15 (m, 4 H, 5-H, 4-H, 3-H), 2.68 (m, 2 H, CH2β(Asn)), 2.40–2.99 (m, 6 H, CH2γ(Pro), CH2γ(Thr), CH2δ(Asn)), 1.60 (m, 2 H, CH2δ(Thr)), 1.44 (dd, J = 7.2 Hz, 3 H, CH3(Asn)), 1.30 (d, J = 7.2 Hz, 3 H, 6-H, 1-H), 1.28–1.17 (m, 12 H, 6 CH2δ(Thr)), 1.21 (d, J = 6.0 Hz, 3 H, CH3(Thr)), 0.87 (m, 3 H, CH3δ(Asn)) ppm. 13C NMR (75 MHz, CDCl3): δC = 173.5, 172.7, 172.4, 171.5, 170.1, 169.7, 167.0, 66.9, 60.2, 55.8, 57.0, 50.1, 47.3, 43.6, 38.5, 36.0, 34.3, 31.4, 29.0, 28.9, 28.8, 28.7, 24.5, 24.3, 22.2, 19.2, 17.1, 16.1, 13.6 ppm. HRMS (EI): calcd. for C31H47N7O8 [M + H]+ 652.4033; found 652.4017.

(3S)-3-[(1R)-1-[(N-Decanoyl-1-prolyl-1-threonyl-1-alanyl-1-L-asparaginyl)amino]ethyl]azetidin-2-one (1b): Decanoyl-PTAN-COOH (4.15 mg, 0.03 mmol) and the aminoazetidinone 9a (3 mg, 0.03 mmol) were dissolved in dry DMF (1.5 mL) and the solution was cooled to 0 °C. HOBT (5.40 mg, 0.04 mmol), EDCI (10.20 mg, 0.06 mmol), and then DIPEA (12 μL, 0.06 mmol) were added to the solution and the reaction was stirred at room temperature for 6 h. The solvent was then evaporated to dryness under vacuum. The residue obtained was dissolved in DCM and washed successively with citric acid solution (1 m) and brine. The aqueous phases were extracted several times with DCM and the resulting organic phases were combined, dried, and concentrated under vacuum. The residue obtained was purified by chromatography on silica gel using DCM/MeOH (9:5.0 to 8:2) as eluent to give 1b (8 mg, 45%) as a white solid. Rf = 0.5 (DCM/MeOH, 9:1). 1H NMR (300 MHz, CDCl3): δH = 5.98 (br., 1 H, NH), 5.12 (quint., J = 6.5 Hz, 1 H, 5-H), 3.42–3.27 (m, 1 H, 3-H), 3.37 (t, J = 5.6 Hz, 1 H, 4-Hb), 3.21 (dd, J = 2.1, J = 5.6 Hz, 4-Ha), 1.43 (d, J = 6.6 Hz, 3 H, 6-H) ppm. 13C NMR (75 MHz, CDCl3): δC = 167.9, 55.2, 54.7, 37.8, 16.2 ppm. HRMS (EI): calcd. for C6H12NO4S [M + H]+ 194.0486; found 194.0494.

(3S)-3-[(1R)-1-[(N-Decanoyl-1-prolyl-1-threonyl-1-alanyl-1-L-asparaginyl)amino]ethyl]azetidin-2-one (1a): Decanoyl-PTAN-COOH (30 mg, 0.03 mmol) and the aminoazetidinone 9b (3 mg, 0.03 mmol) were dissolved in dry DMF (1.50 mL) and the solution was cooled to 0 °C. HOBT (5.40 mg, 0.04 mmol), EDCI (10.20 mg, 0.06 mmol), and then DIPEA (12 μL, 0.06 mmol) were added to the solution and the reaction mixture was stirred at room temperature for 6 h. The solvent was then evaporated to dryness under vacuum. The residue obtained was dissolved in DCM and washed successively with citric acid solution (1 m) and brine. The aqueous phases were extracted several times with DCM and the resulting organic phases were combined, dried, and concentrated under vacuum. The residue obtained was purified by chromatography on silica gel using DCM/MeOH (9:5.0 to 8:2) as eluent to give 1b (8 mg, 45%) as a white solid. Rf = 0.5 (DCM/MeOH, 9:1). 1H NMR (300 MHz, CDCl3): δH = 5.98 (br., 1 H, NH), 5.12 (quint., J = 6.5 Hz, 1 H, 5-H), 3.43 (m, 1 H, 4-H), 3.30–3.24 (m, 2 H, 4-H, 3-H), 1.43 (d, J = 6.6 Hz, 3 H, 6-H) ppm. 13C NMR (75 MHz, CDCl3): δC = 173.5, 172.7, 172.4, 171.5, 170.1, 169.7, 167.0, 66.9, 60.2, 55.8, 57.0, 50.1, 47.3, 43.6, 38.5, 36.0, 34.3, 31.4, 29.0, 28.9, 28.8, 28.7, 24.5, 24.3, 22.2, 19.2, 17.1, 16.1, 13.6 ppm. HRMS (EI): calcd. for C31H47N7O8 [M + H]+ 652.4033; found 652.4017.
during vigorous stirring with 1 M HCl (46 mL, 46 mmol). The phases were separated and the extraction was completed with two additional portions of DCM. The combined organic extracts were washed with water and brine. The organic layer was dried (MgSO4) and concentrated under reduced pressure to afford compound 10 (5.64 g, 70%) as a white solid, which was used in the next step without further purification. An analytical sample was obtained as a white crystalline solid by crystallization from acetone. Rf = 0.5 (DCM/MeOH, 8:2).

1H NMR (300 MHz, CDCl3): δH = 7.92–7.77 (m, 4 H, Hpht), 5.70 and 5.61 (2 × d, J = 3.9 Hz, 2 × 1 H, 5-H, 6-H), 4.71 (s, 1 H, 2-H), 1.85 (s, 3 H, CH3), 1.62 (s, 2 H, CH2) ppm.

13C NMR (75 MHz, CDCl3): δC = 171.1, 168.5, 166.2, 134.2, 131.1, 123.6, 70.6, 66.3, 65.3, 57.9, 30.1, 27.7 ppm. HRMS (EI): calcd. for C16H15N2O5S [M + H]+ 347.0701; found 347.0706.

Additional portions of DCM. The combined organic extracts were dried (MgSO4), and concentrated under vacuum. The residue obtained was purified by chromatography on silica gel using toluene/ethyl acetate (9.5:0.5) as eluent and afforded the desired isomer 11c (2 g, 80% for two steps) as white solids. Rf = 0.3 (toluene/ethyl acetate, 9.5:0.5).

1H NMR (300 MHz, CDCl3): δH = 7.95–7.85 (m, 4 H, Hpht), 7.45–7.34 (m, 5 H, Hpht), 5.57 and 5.45 (2 × d, J = 2.1 Hz, 2 × 1 H, 5-H, 6-H), 5.27 (d, J = 4.2 Hz, 2 H, CH2Ph), 3.93 (s, 1 H, 2-H), 1.67 (s, 3 H, CH3), 1.41 (s, 3 H, CH3) ppm.

13C NMR (75 MHz, CDCl3): δC = 166.2, 166.1, 165.3, 134.5, 134.2, 128.3, 128.3, 128.3, 69.8, 67.5, 66.0, 65.7, 59.5, 31.0, 24.3 ppm. HRMS (EI): calcd. for C23H20N2O5SK [M + K]+ 475.0729; found 475.0708.

Benzyl (2S,5S,6R)-3,3-Dimethyl-7-oxo-6-phenylthiamido-4-thia-azabicyclo[3.2.0]heptane-2-carboxylate (11a): Triethylamine (1.30 mL, 9.39 mmol) was added to benzyl bromide (1.46 mL, 12.20 mmol) to give a mixture of two chlorinated compounds, the desired isomer 11c (2 g, 80% for two steps) as white solids. Rf = 0.3 (toluene/ethyl acetate, 9.5:0.5).

1H NMR (300 MHz, CDCl3): δH = 7.92–7.77 (m, 4 H, Hpht), 7.45–7.34 (m, 5 H, Hpht), 5.57 and 5.45 (2 × d, J = 2.1 Hz, 2 × 1 H, 5-H, 6-H), 5.27 (d, J = 4.2 Hz, 2 H, CH2Ph), 3.93 (s, 1 H, 2-H), 1.67 (s, 3 H, CH3), 1.41 (s, 3 H, CH3) ppm.

13C NMR (75 MHz, CDCl3): δC = 166.2, 166.1, 165.3, 134.5, 134.2, 128.3, 128.3, 69.8, 67.5, 66.0, 65.7, 59.5, 31.0, 24.3 ppm. HRMS (EI): calcd. for C23H20N2O5SK [M + K]+ 475.0729; found 475.0708.

General Procedure 1. Synthesis of Benzoic Acid Derivatives from the Protected Phthalimide Penicillin (Example: Compound 12a): 2-[(2S,5R,6R)-2-(Benzyloxycarbonyl)-3,3-dimethyl-7-oxo-4-thia-1-azabicyclo[3.2.0]heptane-2-carboxylate (12a): Sodium disulfide (0.27 g, 1.14 mmol) was added at 0–5 °C to a solution of the protected phthalimide derivative 11a (0.50 g, 1.14 mmol) in THF/H2O mixture (13 mL/13 mL). The solution was stirred at 0–5 °C for 15 min and then acidified with citric acid (0.5 M) to pH 4. DCM was added and the aqueous phase was extracted three times with DCM. The organic phases were combined and washed successively with water and brine, dried (MgSO4), and concentrated under vacuum. The residue obtained was purified by chromatography on silica gel using DCM/ethyl acetate (9.5:0.5) as eluent and afforded the desired isomer 12a (0.35 g, 67%) as a white solid. Rf = 0.2 (DCM/ethyl acetate, 4:6).

1H NMR (300 MHz, CDCl3): δH = 7.92–7.77 (m, 4 H, Hpht), 7.45–7.34 (m, 5 H, Hpht), 5.57 and 5.45 (2 × d, J = 2.1 Hz, 2 × 1 H, 5-H, 6-H), 5.27 (d, J = 4.2 Hz, 2 H, CH2Ph), 3.93 (s, 1 H, 2-H), 1.67 (s, 3 H, CH3), 1.41 (s, 3 H, CH3) ppm.
General Procedure 2. Synthesis of the Phthalosimido Derivative from the Benzozio Acid Derivative (Example: Compound 13a)  

**Benzyl (2S,5S,6R)-3-Dimethyl-7-oxo-6-phthalisimido-4-thia-1-azabicyclo[3.2.0]heptane-2-carboxylate (13a):** DCC (0.12 g, 0.60 mmol) was added to a stirred solution of the acid 12a (0.27 g, 0.60 mmol) in dry DCM (10 mL) at 0 °C. The reaction mixture was stirred for 1 h at 0 °C and for an additional 1 h at room temperature. Then the reaction mixture was again cooled to 0 °C and the precipitated dicyclohexylurea (DCU) was removed by filtration. The residue obtained was purified by chromatography on silica gel using DCM/ethyl acetate (9:1) as eluent to give 13a (0.18 g, 70%) as a white foam. Rₜ = 0.5 (hexane/ethyl acetate, 7:4). ¹H NMR (300 MHz, CDCl₃): δₗ = 8.04 (d, J = 7.4 Hz, 1 H, H₂pht), 7.96 (d, J = 7.2 Hz, 1 H, H₁pht), 7.83 (td, J = 7.2, J = 1.2 Hz, 1 H, H₁pht), 7.76 (td, J = 7.4, J = 1.2 Hz, 1 H, H₁pht), 7.42–7.35 (m, 5 H, H₂pht), 5.71 and 5.63 (2 x d, J = 4.2 Hz, 2 x 1 H, H₅, H₆), 5.23 (d, J = 4.2 Hz, 1 H, CH₂Ph), 4.55 (s, 1 H, H₂pht), 1.66 (s, 3 H, CH₃), 1.43 (s, 3 H, CH₂pht). ¹³C NMR (75 MHz, CDCl₃): δC = 172.2, 172.4, 163.2, 161.5, 151.3, 153.2, 153.4, 133.1, 128.7, 127.5, 121.5, 123.5, 70.4, 68.0, 67.3, 67.1, 63.3, 30.6, 26.2 ppm. HRMS (EI): calcd. for C₂₃H₂₃N₂O₅Na [M + Na⁺] 459.0990; found 459.0983.

**Benzyl (2S,5S,6S)-3-Dimethyl-7-oxo-6-phthalisimido-4-thia-1-azabicyclo[3.2.0]heptane-2-carboxylate (13b):** General procedure 2 was carried out by using compound 12b (0.14 g, 0.32 mmol) as the starting material. The residue obtained was purified by chromatography on silica gel using DCM/ethyl acetate (9:1) as eluent to give 13b (0.07 g, 57%) as a white foam. Rₜ = 0.4 (hexane/ethyl acetate, 7:4). ¹H NMR (300 MHz, CDCl₃): δₗ = 7.98 (t, J = 6.5 Hz, 2 H, H₂pht), 7.84 (dt, J = 7.2, J = 1.2 Hz, 1 H, H₁pht), 7.70 (dt, J = 7.2, J = 0.9 Hz, 1 H, H₁pht), 7.42–7.36 (m, 5 H, H₂pht), 5.53 and 5.40 (2 x d, J = 1.5 Hz, 2 x 1 H, 5-H, 6-H), 5.23 (d, J = 4.8 Hz, 2 H, CH₂Ph), 4.62 (s, 1 H, 2-H), 1.66 (s, 3 H, CH₃), 1.44 (s, 3 H, CH₂pht). ¹³C NMR (75 MHz, CDCl₃): δC = 168.3, 167.0, 163.3, 151.4, 153.5, 153.4, 133.1, 128.4, 128.3, 125.2, 123.3, 74.6, 69.5, 69.3, 67.0, 64.3, 33.2, 25.5 ppm. HRMS (EI): calcd. for C₂₃H₂₃N₂O₅Na [M + Na⁺] 459.0990; found 459.0989.

**Benzyl (2S,5S,6S)-3-Dimethyl-7-oxo-6-phthalisimido-4-thia-1-azabicyclo[3.2.0]heptane-2-carboxylate (13c):** General procedure 2 was carried out by using compound 12c (0.30 g, 0.66 mmol) as the starting material. The residue obtained was purified by chromatography on silica gel using DCM/ethyl acetate (9:1) as eluent to give 13c (0.12 g, 45%) as a white foam. Rₜ = 0.4 (hexane/ethyl acetate, 7:4). ¹H NMR (300 MHz, CDCl₃): δₗ = 7.98 (t, J = 6.5 Hz, 2 H, H₂pht), 7.84 (td, J = 7.2, J = 1.2 Hz, 1 H, H₁pht), 7.70 (td, J = 7.2, J = 0.9 Hz, 1 H, H₁pht), 7.42–7.36 (m, 5 H, H₂pht), 5.53 and 5.40 (2 x d, J = 1.5 Hz, 2 x 1 H, 5-H, 6-H), 5.23 (d, J = 4.8 Hz, 2 H, CH₂Ph), 4.62 (s, 1 H, 2-H), 1.66 (s, 3 H, CH₃), 1.44 (s, 3 H, CH₂pht). ¹³C NMR (75 MHz, CDCl₃): δC = 166.6, 165.8, 163.2, 151.5, 153.5, 153.4, 133.1, 128.6, 128.3, 128.0, 125.2, 123.3, 71.3, 70.4, 67.5, 67.0, 64.8, 30.3, 24.9 ppm. HRMS (EI): calcd. for C₂₃H₂₃N₂O₅Na [M + Na⁺] 459.0990; found 459.0996.
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fication because of the total degradation of the compound on silica gel chromatography. $R_f = 0.4$ (DCM/ethyl acetate, 9:8:0.2). Major product $13d$: $^1$H NMR (300 MHz, CDCl$_3$): $\delta_{H}$ = 8.04 (d, $J = 7.5$ Hz, 1 H, $H_{ph}$), 7.95 (d, $J = 7.5$ Hz, 1 H, $H_{ph}$), 7.85–7.73 (m, 2 H, $H_{ph}$), 7.44–7.34 (m, 5 H, $H_{Ph}$), 5.52 and 5.46 (2 × d, $J = 4.1$ Hz, 2 × 1 H, 6-H, 5-H), 5.27 (s, 2 H, $CH_{2}Ph$), 3.91 (s, 1 H, 2-H), 1.64 (s, 3 H, $CH_{3}$), 1.54 (s, 3 H, $CH_{3}$) ppm. HRMS (EI): calcd. for $C_{15}H_{20}N_{2}O_{5}$SnNa [M + Na]$^+$ 459.1002; found 459.0985.

General Procedure 3. Deprotection of the Phthalisomide Group (Example: Compound $14a$)

Benzyl (2S,5R,6R)-6-Amino-3,3-dimethyl-7-oxo-4-thia-1-azabicyclo[3.2.0]heptane-2-carboxylate ($14a$): The phthalisomide $13a$ (0.17 g, 0.38 mmol) was dissolved in dry THF (8.50 mL) and the solution was cooled to −20 °C. A 2 m solution of $N$-methylhydrazine (0.19 mL, 0.38 mmol) in THF was added dropwise and the reaction mixture was stirred for 20 min at −20 °C. The solvent was evaporated under reduced pressure and the residue obtained was co-evaporated several times with chloroform until precipitation of the byproduct hydrazide. The byproduct was removed by filtration and the filtrate was concentrated to afford the crude amine (0.19 mL, 0.38 mmol) as a white amorphous solid. No further purification because of the total degradation of the compound on silica gel chromatography. $R_f = 0.4$ (DCM/ethyl acetate, 6:4). $\delta_{H}$ = 8.04 (d, $J = 3.5$ Hz, 1 H, $H_{ph}$), 7.95 (d, $J = 7.5$ Hz, 1 H, $H_{ph}$), 7.85–7.73 (m, 2 H, $H_{ph}$), 7.44–7.34 (m, 5 H, $H_{Ph}$), 5.52 and 5.46 (2 × d, $J = 4.1$ Hz, 2 × 1 H, 6-H, 5-H), 5.27 (s, 2 H, $CH_{2}Ph$), 3.91 (s, 1 H, 2-H), 1.64 (s, 3 H, $CH_{3}$), 1.54 (s, 3 H, $CH_{3}$) ppm. HRMS (EI): calcd. for $C_{15}H_{20}N_{2}O_{5}$SnNa [M + Na]$^+$ 459.0999; found 459.0985.

Benzyl (2S,5R,6R)-6-Amino-3,3-dimethyl-7-oxo-4-thia-1-azabicyclo[3.2.0]heptane-2-carboxylate ($14a$): The phthalisomide $13a$ (0.17 g, 0.38 mmol) was dissolved in dry THF (8.50 mL) and the solution was cooled to −20 °C. A 2 m solution of $N$-methylhydrazine (0.19 mL, 0.38 mmol) in THF was added dropwise and the reaction mixture was stirred for 20 min at −20 °C. The solvent was evaporated under reduced pressure and the residue obtained was co-evaporated several times with chloroform until precipitation of the byproduct hydrazide. The byproduct was removed by filtration and the filtrate was concentrated to afford the crude amine (0.19 mL, 0.38 mmol) as a white amorphous solid. No further purification because of the total degradation of the compound on silica gel chromatography. $R_f = 0.4$ (DCM/ethyl acetate, 6:4). $\delta_{H}$ = 8.04 (d, $J = 3.5$ Hz, 1 H, $H_{ph}$), 7.95 (d, $J = 7.5$ Hz, 1 H, $H_{ph}$), 7.85–7.73 (m, 2 H, $H_{ph}$), 7.44–7.34 (m, 5 H, $H_{Ph}$), 5.52 and 5.46 (2 × d, $J = 4.1$ Hz, 2 × 1 H, 6-H, 5-H), 5.27 (s, 2 H, $CH_{2}Ph$), 3.91 (s, 1 H, 2-H), 1.64 (s, 3 H, $CH_{3}$), 1.54 (s, 3 H, $CH_{3}$) ppm. HRMS (EI): calcd. for $C_{15}H_{20}N_{2}O_{5}$SnNa [M + Na]$^+$ 459.0999; found 459.0985.

Benzyl (2S,5R,6R)-6-Amino-3,3-dimethyl-7-oxo-4-thia-1-azabicyclo[3.2.0]heptane-2-carboxylate ($14a$): The phthalisomide $13a$ (0.17 g, 0.38 mmol) was dissolved in dry THF (8.50 mL) and the solution was cooled to −20 °C. A 2 m solution of $N$-methylhydrazine (0.19 mL, 0.38 mmol) in THF was added dropwise and the reaction mixture was stirred for 20 min at −20 °C. The solvent was evaporated under reduced pressure and the residue obtained was co-evaporated several times with chloroform until precipitation of the byproduct hydrazide. The byproduct was removed by filtration and the filtrate was concentrated to afford the crude amine (0.19 mL, 0.38 mmol) as a white amorphous solid. No further purification because of the total degradation of the compound on silica gel chromatography. $R_f = 0.4$ (DCM/ethyl acetate, 6:4). $\delta_{H}$ = 8.04 (d, $J = 3.5$ Hz, 1 H, $H_{ph}$), 7.95 (d, $J = 7.5$ Hz, 1 H, $H_{ph}$), 7.85–7.73 (m, 2 H, $H_{ph}$), 7.44–7.34 (m, 5 H, $H_{Ph}$), 5.52 and 5.46 (2 × d, $J = 4.1$ Hz, 2 × 1 H, 6-H, 5-H), 5.27 (s, 2 H, $CH_{2}Ph$), 3.91 (s, 1 H, 2-H), 1.64 (s, 3 H, $CH_{3}$), 1.54 (s, 3 H, $CH_{3}$) ppm. HRMS (EI): calcd. for $C_{15}H_{20}N_{2}O_{5}$SnNa [M + Na]$^+$ 459.0999; found 459.0985.

General Procedure 3. Deprotection of the Phthalisomide Group (Example: Compound $14a$)

Benzyl (2S,5R,6R)-6-Amino-3,3-dimethyl-7-oxo-4-thia-1-azabicyclo[3.2.0]heptane-2-carboxylate ($14a$): The phthalisomide $13a$ (0.17 g, 0.38 mmol) was dissolved in dry THF (8.50 mL) and the solution was cooled to −20 °C. A 2 m solution of $N$-methylhydrazine (0.19 mL, 0.38 mmol) in THF was added dropwise and the reaction mixture was stirred for 20 min at −20 °C. The solvent was evaporated under reduced pressure and the residue obtained was co-evaporated several times with chloroform until precipitation of the byproduct hydrazide. The byproduct was removed by filtration and the filtrate was concentrated to afford the crude amine (0.19 mL, 0.38 mmol) as a white amorphous solid. No further purification because of the total degradation of the compound on silica gel chromatography. $R_f = 0.4$ (DCM/ethyl acetate, 6:4). $\delta_{H}$ = 8.04 (d, $J = 3.5$ Hz, 1 H, $H_{ph}$), 7.95 (d, $J = 7.5$ Hz, 1 H, $H_{ph}$), 7.85–7.73 (m, 2 H, $H_{ph}$), 7.44–7.34 (m, 5 H, $H_{Ph}$), 5.52 and 5.46 (2 × d, $J = 4.1$ Hz, 2 × 1 H, 6-H, 5-H), 5.27 (s, 2 H, $CH_{2}Ph$), 3.91 (s, 1 H, 2-H), 1.64 (s, 3 H, $CH_{3}$), 1.54 (s, 3 H, $CH_{3}$) ppm. HRMS (EI): calcd. for $C_{15}H_{20}N_{2}O_{5}$SnNa [M + Na]$^+$ 459.0999; found 459.0985.
Benzyl (2S,5R,6S)-6-[(N-Decanoyl-t-prolyl-t-threonyl-l-alanyl-l-threonyl-l-asparaginyl)amino]-3,3-dimethyl-7-oxo-4-thia-1-azabicyclo[3.2.0]-heptane-2-carboxylate (2b): Decanoyl-PTAN-COOH (4: 20 mg, 0.03 mmol) and the amine 14b (16.50 mg, 0.05 mmol) were dissolved in dry DMF (1.5 mL) and the solution was cooled to 0 °C. HOBr (7.20 mg, 0.05 mmol) and EDCI (13.80 mg, 0.06 mmol) were added to the solution and the reaction mixture was stirred at room temperature for 6 h. The solvent was then evaporated to dryness under vacuum. The residue obtained was dissolved in DCM and washed successively with citric acid solution (1%), water, saturated NaHCO₃ solution, and brine. The aqueous phases were extracted several times with DCM and the resulting organic phases were combined, dried, and concentrated under vacuum. The residue obtained was purified by chromatography on silica gel using DCM/MeOH (9:5.0:5 to 9:1) as a gradient to give 2b (14 mg, 46%) as a white solid. Rf = 0.3 (DCM/MeOH, 9:1). ¹H NMR (300 MHz, CDCl₃/CD3OD): δH = 8.28, 7.87, 7.82, 6.75, 6.24 (5 × br., 5 × 1 H, 4NHe, OHe), 7.37 (m, 5 H, HPh), 5.28 (d, J = 1.2 Hz, 1 H, 5-H), 5.18 (s, 2 H, CH₂Ph), 5.02 (dd, J = 1.2, J = 7.8 Hz, 1 H, 6-H), 4.83 (m, 1 H, Hα(Asn)), 4.51–4.27 (m, 4 H, Hα(Ala, Pro, Thr), Hβ(Thr)), 3.69, 3.52 (m, 2 H, CH₂β(Pro)), 2.87 (dd, J = 15.5, J = 6.6 Hz, 1 H, CH₂β(Asn)), 2.70 (dd, J = 15.5, J = 5.1 Hz, 1 H, CH₂β(β(Asn)), 2.38–2.29 (m, 2 H, CH₂β(Pro)), 2.25–1.97 (m, 4 H, CH₂β(Pro), 1.61 (m, 2 H, CH₂β(Asn)), 1.56 (s, 3 H, CH₃), 1.44 (d, J = 7.3 Hz, 3 H, CH₃(Ala), 1.36 (s, 3 H, CH₃), 1.31–1.25 (m, 12 H, 6 CH₃deca), 1.17 (d, J = 6.3 Hz, 3 H, CH₃(CH₃)), 0.88 (t, J = 6.9 Hz, 3 H, CH₃deca) ppm. ¹³C NMR (75 MHz, CDCl₃/CD3OD): δC = 173.2, 172.9, 172.5, 172.4, 170.6, 169.9, 167.1, 134.5, 128.9, 69.4, 68.8, 67.0, 66.7, 66.3, 60.8, 53.2, 50.5, 49.7, 47.4, 36.3, 34.4, 33.0, 31.5, 29.3, 29.2, 29.1, 28.9, 25.4, 24.7, 24.4, 22.3, 19.3, 16.4, 13.8 ppm. HRMS (EI): calcd. for C₄₅H₆₃N₇O₁₀SNa [M + Na]⁺ 866.4098; found 866.4094.

Benzyl (2S,5S,6R)-6-[(N-Decanoyl-t-prolyl-t-threonyl-l-alanyl-l-threonyl-l-asparaginyl)amino]-3,3-dimethyl-7-oxo-4-thia-1-azabicyclo[3.2.0]-heptane-2-carboxylate (2c): Decanoyl-PTAN-COOH (4: 25 mg, 0.04 mmol) and the amine 14c (27 mg, 0.09 mmol) were dissolved in dry DMF (2 mL) and the solution was cooled to 0 °C. HOBr (9.1 mg, 0.05 mmol) and EDCI (17 mg, 0.09 mmol) were added to the solution and the reaction mixture was stirred at room temperature for 6 h. The solvent was then evaporated to dryness under vacuum. The residue obtained was dissolved in DCM and washed successively with water, saturated NaHCO₃ solution, and brine. The aqueous phases were extracted several times with DCM and the resulting organic phases were combined, dried, and concentrated under vacuum. The residue obtained was purified by chromatography on silica gel using DCM/IPrOH (9:5.0:5 to 8:2) as a gradient to give 2c (19 mg, 68%) as a white solid. Rf = 0.3 (DCM/IPrOH, 9:1). ¹H NMR (300 MHz, CDCl₃/CD3OD): δH = 8.04, 7.50, 7.45, 6.54, 6.15 (5 × br., 5 × 1 H, 4NHe, OHe), 7.39–7.32 (m, 5 H, HPh), 5.41 (dd, J = 4.1, J = 7.8 Hz, 1 H, 6-H), 5.23 (d, J = 4.1 Hz, 1 H, 5-H), 5.22 (d, J = 3.6 Hz, 2 H, CH₂Ph), 4.88 (m, 1 H, Hα(Asn)), 4.52–4.32 (m, 4 H, Hα(Ala, Pro, Thr), Hβ(Thr)), 3.84 (s, J = 2.0 Hz), 3.63, 3.48 (m, 2 H, CH₂β(Pro)), 2.82 (dd, J = 15.5, J = 6 Hz, 1 H, CH₂β(β(Asn)), 2.71 (dd, J = 15.5, J = 5.5 Hz, 1 H, CH₂β(β(Asn)), 2.34–2.28 (m, 2 H, CH₂β(Pro)), 2.15–1.80 (m, 4 H, CH₂β(Pro), CH₂deca)), 1.62 (s, 3 H, CH₃), 1.59 (m, 2 H, CH₂deca), 1.48 (3 H, CH₃), 1.43 (d, J = 7.0 Hz, 3 H, CH₃(Ala)), 1.30–1.22 (m, 12 H, 6 CH₃deca), 1.16 (d, J = 6.0 Hz, 3 H, CH₃(CH₃)), 0.87 (t, J = 7.0 Hz, 3 H, 3 CH₃deca) ppm. ¹⁴C NMR (75 MHz, CDCl₃/CD3OD): δC = 173.2, 172.9, 172.5, 172.4, 170.6, 170.5, 170.4, 165.1, 134.4, 128.3, 71.4, 67.3, 66.9, 63.8, 62.6, 60.2, 59.1, 58.0, 49.6, 49.5, 47.4, 36.5, 34.4, 31.5, 29.3, 29.1, 29.0, 28.9, 28.7, 26.6, 24.7, 24.4, 22.3, 19.0, 17.1, 13.7 ppm. HRMS (EI): calcd. for C₄₅H₆₁N₇O₁₀SNa [M + Na]⁺ 866.4109; found 866.4109.

Supporting Information (see footnote on the first page of this article): Experimental characterization data (¹H and ¹³C NMR, HRMS) for all the compounds synthesized and procedures for the biological in vitro evaluation.

Acknowledgments

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\textbf{β-Lactam Lipopeptides as Bacterial Signal Peptidase I Inhibitors}


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We present the design, asymmetric synthesis, and biological evaluation of novel β-lactam lipopeptides, inhibitors of the signal peptidase I, an essential bacterial enzyme for the viability of both Gram-negative and Gram-positive bacteria. The importance of the azetidinone moiety is demonstrated and can serve as a starting point to improve the structure of these new kinds of SPase I inhibitors.