Solid-Phase Synthesis of Biaryl Cyclic Lipopeptides Derived from Arylomycins

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ABSTRACT: An efficient approach for the solid-phase synthesis of N-methylated tailed biaryl cyclic lipopeptides based on the structure of arylomycins was established. Each of these analogues incorporates an N-terminal linear lipopeptide attached to a biaryl cyclic tripeptide containing a Phe−Tyr, a Tyr−Tyr, or a His−Tyr linkage. This methodology first involved an intramolecular Suzuki−Miyaura arylation of a linear peptidyl resin incorporating the corresponding halogenated amino acid at the N-terminus and a boronotyrosine at the C-terminus. After N-methylation of the resulting biaryl cyclic peptidyl resin, the N-methylated lipopeptidyl tail was then assembled. The biaryl cyclic lipopeptides were purified and characterized.

■ INTRODUCTION

Biaryl cyclic peptides have been shown to provide a promising scaffold for the development of novel drugs, owing to their target selectivity, binding affinity, and low toxicity. Thus, they have attracted attention from fields including medicinal chemistry and chemical biology. 1-3 The aryl−aryl bonds in these compounds commonly involve the linkage between the side chains of two aromatic amino acids. Among natural biaryl cyclic peptides, arylomycins A and B are a class of biaryl-containing peptide antibiotics that contain a lipopeptidyl tail attached to a biaryl cyclic tripeptide core. 4-6 The peptidyl tail is a tripeptide with the N-terminus methylated and acylated with a fatty acid of 12−16 carbon atoms. The cyclic core contains an N-methylated residue and incorporates a biaryl linkage between a 4-hydroxy-L-phenylglycine derivative and a tyrosine residue (Figure 1). In particular, the A series of arylomycins possesses an unmodified core, while the compounds of the B series have a nitro substituent on the phenol ring of tyrosine. Arylomycins were isolated from the fermentation broth of Streptomyces sp. Tü 6075 and display significant activity against a variety of Gram-positive and Gram-negative bacteria that results from the inhibition of bacterial type I signal peptidase (SPase). In addition, it has been shown that, even at sub-minimal inhibitory concentration (MIC) levels, arylomycins are able to inhibit a range of virulence processes as well as the dissemination of resistance. 1-4,12

The structural features of arylomycins in conjunction with their significant biological activities have prompted scientists to...
develop efficient methods to obtain these compounds and a diversity of analogues.\textsuperscript{4,6,13–21} In particular, Romesberg and co-workers reported the total synthesis of arylomycin A\textsubscript{2} in solution.\textsuperscript{4} They proposed two strategies that differed in the macrocyclization step for the formation of the biaryl macrocyclic tripeptide core. This step was performed by standard macrolactamization of a conveniently protected biaryl linear tripeptide. Alternatively, the cyclization was carried out through an intramolecular Suzuki−Miyaura cross-coupling of a tripeptide containing a 3-iodo-L-phenylglycine and a 3-borono-L-tyrosine. The latter macrocyclization strategy resulted in being the most successful. The lipopeptidyl tail was then assembled to the biaryl core providing arylomycin A\textsubscript{2}. A similar strategy was reported by Zhu and co-workers for the total synthesis of arylomycins A\textsubscript{2} and B\textsubscript{2} via an intramolecular Suzuki−Miyaura reaction.\textsuperscript{14} These methodologies were extended to the preparation of other arylomycins \textsuperscript{5,6} and derivatives.\textsuperscript{15,16} Recent reports have been focused on optimizing the synthesis of the macrocyclic core of arylomycins. To this end, Romesberg and Baran have described the scalable macrocyclization of the Hpg−Ala−Tyr precursor, using a stoichiometric Cu-mediated oxidative coupling.\textsuperscript{17} Lim and co-workers have developed protocols for the preparation of the macrocyclic core of arylomycins using amide bond formation or a Suzuki−Miyaura cross-coupling for the macrocyclization step.\textsuperscript{19,20} More recently, the Pappo group has devised a three-step synthesis for the cyclic core involving an activating-group-assisted catalytic oxidative coupling for the formation of the biaryl bond followed by macrolactamization.\textsuperscript{21}

Since conventional solution-phase synthesis requires very long and tedious work-up procedures and troublesome purifications after each reaction step, the solid-phase peptide synthesis has become a more powerful method for accelerating such a process.\textsuperscript{22} Moreover, solid-supported chemistry has the additional advantage of allowing the preparation of analogues in a convenient and flexible manner. However, to the best of our knowledge, the solid-phase synthesis of arylomycins has not yet been reported. In fact, the preparation of biaryl peptides on a solid support has been scarcely developed.\textsuperscript{23–28} In this context and within our interest in the synthesis of biaryl peptides, since 2008 we have established methodologies for the preparation of biaryl cyclic peptides centered on the formation of the aryl–aryl bond through a solid-phase Suzuki−Miyaura cross-coupling.\textsuperscript{29} We chose this reaction for the formation of the biaryl bond because of its widely reported advantages. In particular, we have described the synthesis of linear and cyclic peptides containing a biaryl bond between the side chains of a Phe, a Tyr, or a His residue.\textsuperscript{30–36} Based on the interest arisen from arylomycins, we decided to extend our solid-phase methodology to the preparation of biaryl cyclic lipopeptides related to arylomycins. In particular, we focused our attention on analogues 1−3, which contain a Phe−Tyr, a Tyr−Tyr, or a His−Tyr biaryl linkage, respectively (Figure 2).

\section*{RESULTS AND DISCUSSION}

\textbf{Retrosynthetic Analysis.} The solid-phase synthesis of the biaryl cyclic lipopeptides 1−3 was planned according to the retrosynthetic analysis depicted in Scheme 1. It would involve the synthesis of the biaryl cyclic peptidyl resins 4−6 through intramolecular Suzuki−Miyaura cross-coupling of the tripeptidyl resins 7−9, incorporating a borotyrosine and a halogenated aromatic amino acid. Subsequent N-methylation of 4−6, elongation of the peptidyl tail, methylation of the N-
Synthesis of the Biaryl Cyclic Peptidyl Resins 4–6

According to the retrosynthetic analysis, the preparation of biaryl cyclic lipopeptides 1–3 required the synthesis of biaryl cyclic peptidyl resins 4–6 (Scheme 1). We first investigated the preparation of 4, bearing a Phe–Tyr linkage, from the tripeptidyl resin Boc-Phe(4-I)-Ala-Tyr(3-BPin,Me)-Rink-MBHA (7), which incorporates a 3-boronotyrosine and a 4-iodophenylalanine (Scheme 2). The synthesis of 7 started from an Fmoc-Rink-MBHA resin (0.4 mmol/g) and followed a standard 9-fluorenylmethoxycarbonyl (Fmoc)/tert-butyl (tBu) strategy. First, the dipeptidyl resin Fmoc-Ala-Tyr(3-I,Me)-Rink-MBHA (10) was obtained through sequential Fmoc removal and coupling steps. The Fmoc group was removed by treatment with piperidine/dimethylformamide (DMF) (3:7); coupling of Fmoc-Tyr(3-I,Me)-OH was mediated by 1-[(1-(cyano-2-ethoxy-2-oxo ethylidine aminoxy)dimethylaminomorpholino)] uronium hexafluorophosphate (COMU), ethyl 2-cyano-2-(hydroxyimino)acetate (Oxyma), and N,N-diisopropylethylamine (DIEA) in DMF; and Fmoc-Ala-OH was coupled using N,N′-diisopropylcarbodiimide (DIPCDI) and Oxyma in DMF. An aliquot of resin 10 was treated with trifluoroacetic acid (TFA)/H2O/triisopropylsilane (TIS) (95:2.5:2.5) for 2 h at room temperature, providing Fmoc-Ala-Tyr(3-I,Me)-NH2 in 98% purity. Then, the Fmoc group of 10 was replaced by a trityl (Tr) group due to the lability of the former to the basic reaction conditions of the next borylation step. After deprotection of 10, tritylation of the resulting resin was performed with trityl chloride (TrCl) and DIEA in DMF, leading to resin Tr-Ala-Tyr(3-I,Me)-Rink-MBHA (11). Acidolytic cleavage of an aliquot of this resin gave H-Ala-Tyr(3-I,Me)-NH2 in 92% purity. Next, 11 was subjected to Miyaura borylation by exposure to the conditions previously described in our group, which involve the use of bis(pinacolato)diboron (B2Pin2) (4 equiv), PdCl2(dppf) (0.18 equiv), 1,1′-bis(diphenylphosphanyl)ferrocene (dppf) (0.09 equiv), and KOAc (6 equiv) in anhydrous dimethyl sulfoxide (DMSO) at 80 °C for 8 h. We observed that longer reaction times led to the protodeborylation of the corresponding linear precursor. Acidolytic cleavage of an aliquot of the resulting resin Tr-Ala-Tyr(3-BPin,Me)-Rink-MBHA (12) yielded the expected borono peptide H-Ala-Tyr(3-B(OH)2,Me)-NH2 in 72% purity. This boronic acid resulted from the hydrolysis of the pinacol boronic ester during high-performance liquid chromatography (HPLC) analysis, which was confirmed by mass spectrometry. Next, selective trityl group removal of 12 by treatment with TFA/H2O/CH2Cl2 (0.2:1:98.8), followed by coupling of Boc-Phe(4-I)-OH using DIPCDI and Oxyma in DMF for 3 h, provided the tripeptidyl resin Boc-Phe(4-I)-Ala-Tyr(3-B(OH)2,Me)-Rink-MBHA (7). An aliquot of this resin was acidolytically cleaved, affording H-Phe(4-I)-Ala-Tyr(3-B(OH)2,Me)-NH2 in 81% purity, which was characterized by mass spectrometry.

Based on our previous experience on the synthesis of biaryl cyclic peptides containing a Phe–Tyr linkage, macrocyclization of the linear peptidyl resin 7 was achieved via a Suzuki–Miyaura cross-coupling using Pd2(dba)3 (0.2 equiv) as the catalyst, SPhos (0.4 equiv) as the ligand, and KF (4 equiv) as the base in degassed 1,2-dimethoxyethane (DME)/EtOH/
H$_2$O (9:9:2) under microwave irradiation at 120 °C for 30 min (Scheme 2). Giralt and co-workers reported that the presence of Buchwald's SPhos ligand avoids racemization of α-amino acids, such as tyrosine derivatives in Suzuki–Miyaura reactions. An aliquot of the resulting resin 4 was cleaved, and HPLC and mass spectrometry analysis of the crude reaction mixture revealed the formation of the expected biaryl cyclic tripeptide 13 in 92% purity together with traces of the protodeborylated byproduct H-Phe(4-I)-Ala-Tyr(Me)-NH$_2$, which is a common byproduct in Suzuki–Miyaura reactions. Biaryl cyclic tripeptide 13 was isolated by reverse-phase column chromatography and analyzed by HPLC and was obtained in 99% purity. It was also characterized by mass spectrometry and nuclear magnetic resonance (NMR) spectroscopy.

The strategy described for resin 4 was then applied to the synthesis of the biaryl cyclic peptidyl resin 5, bearing a Tyr−Tyr linkage (Scheme 2). Accordingly, the tripeptidyl resin Boc-Tyr(3-I,Me)-Ala-Tyr(3-BPin,Me)-Rink-MBHA (8), containing a 3-boronotyrosine and a 3-iodotyrosine, was prepared from Tr-Ala-Tyr(3-BPin,Me)-Rink-MBHA (12) by trityl group removal and coupling of Boc-Tyr(3-I,Me)-OH mediated by DIPCDI and Oxyma in DMF. Acidolytic cleavage of an aliquot of 8 afforded H-Tyr(3-I,Me)-Ala-Tyr(3-B(OH)$_2$,Me)-NH$_2$ in 82% purity, which was characterized by mass spectrometry. Resin 8 was then subjected to Suzuki–Miyaura macrocyclization under the same conditions employed for the cyclization of resin 7. HPLC and mass spectrometry analysis of the crude reaction mixture from the cleavage of an aliquot of the resulting resin 5 revealed the formation of the expected biaryl cyclic tripeptide 14 in 73% purity. Similar to 13, this biaryl cyclic peptide 14 was obtained together with traces of the protodeborylated byproduct H-Tyr(3-I,Me)-Ala-Tyr(Me)-NH$_2$. The biaryl cyclic peptide 14 was purified by column chromatography. It was obtained in >99% HPLC purity and characterized by mass spectrometry and NMR spectroscopy.

Similarly, we attempted the synthesis of resins 6 bearing a His−Tyr linkage (Scheme 2). In this case, after trityl group removal of Tr-Ala-Tyr(3-BPin,Me)-Rink-MBHA (12), the regioisomeric mixture of Boc-His(5-Br,1-SEM)-OH and Boc-His(5-Br,3-SEM)-OH was coupled to the free amino group, affording resins Boc-His(5-Br,1-SEM)-Ala-Tyr(3-BPin,Me)-Rink-MBHA (9a) and Boc-His(5-Br,3-SEM)-Ala-Tyr(3-BPin,Me)-Rink-MBHA (9b). TFA/H$_2$O/TIS treatment of an aliquot of 9 under stirring for 3 h yielded H-His(5-Br)-Ala-Tyr(3-B(OH)$_2$,Me)-NH$_2$ in 82% purity, and it was characterized by mass spectrometry. Subsequent intramolecular Suzuki–Miyaura macrocyclization of resins 9 at 140 °C provided the expected regioisomeric biaryl cyclic tripeptidyl resins 6. An aliquot of these resins was cleaved and analyzed by HPLC and mass spectrometry. We observed the formation of the expected biaryl cyclic tripeptide 15 in 65% purity together with the protodeborylated H-His(5-Br)-Ala-Tyr(3-B(OH)$_2$,Me)-NH$_2$ and oxidized H-His(5-Br)-Ala-Tyr(3-OH,Me)-NH$_2$ byproducts, with the latter also being a common side product from Suzuki–Miyaura cross-couplings. In agreement with previous reports, this result revealed that the macrocyclization through the formation of a His−Tyr linkage is more difficult than that involving a Phe−Tyr or a Tyr−Tyr bond. The biaryl cyclic tripeptide 15 was purified by column chromatography (75% HPLC purity) and characterized by mass spectrometry and NMR spectroscopy.
Scheme 4. Synthesis of Biaryl Cyclic Lipopeptides 31–33

Synthesis of the Biaryl Cyclic Lipohexapeptides 1–3. With the peptidyl resins 4–6 in hand, we then proceeded to the N-methylation of the biaryl unit and the elongation of the lipopeptidyl tail to obtain the biaryl cyclic lipohexapeptides 1–3 (Scheme 3). For this purpose, the N-terminal Boc group of peptidyl resins 4–6 was selectively removed under mild conditions, using trimethylsilyl triflate (TMSOTf) in the presence of 2,6-lutidine in CH$_2$Cl$_2$. N-methylation was then performed following a three-step procedure that involved protection of the N-terminal amine with an o-nitrobenzenesulfonyl (oNBS) group, followed by Mitsunobu N-methylation of the sulfonamide and oNBS group removal. Protection of the free amino group was achieved with oNBS-Cl and DIEA in CH$_2$Cl$_2$. Mitsunobu N-methylation required the use of PPh$_3$ and DIPCDI and Oxyma in DMF. The completion of the reaction was checked with the chloranil test. The resulting biaryl cyclic lipopeptidyl resins were acidolytically cleaved and analyzed by HPLC and mass spectrometry. However, only traces of the expected palmitoylated sequences 1–3 were obtained, with the major product peptides being 25–27 without the palmitoyl group. Although different assays were performed, we were not able to improve the acylation of peptidyl resins 22–24.

Despite the above results, we decided to assay the incorporation of an amino acid at the N-terminus of the peptidyl tail of resins 22–24 instead of palmitic acid. A Lys residue was selected because it could favor the analysis of the biaryl cyclic lipopeptides by mass spectrometry (Scheme 4). Thus, Fmoc-Lys(Boc)-OH was coupled to the N-terminus of the tripeptidyl tail of resins using DIPCDI and Oxyma in DMF overnight. After Fmoc group removal, they were acylated with palmitic acid (Scheme 3). Thus, Fmoc-Gly-OH, Fmoc-Lys(Boc)-OH, and Fmoc-D-Ser(Bu)-OH were employed as Fmoc-protected amino acids and were incorporated into the resin using DIPCDI and Oxyma in DMF. The completion of the couplings onto primary and secondary amines was monitored with the ninhydrin and chloranil tests, respectively. An aliquot of the resulting tripeptidyl tail of resins 19–21 was cleaved under acyclic conditions, providing the corresponding biaryl cyclic hexapeptides bearing a Phe–Tyr, a Tyr–Tyr, and a His–Tyr linkage in 85, 75, and 18% HPLC purity, respectively, which were characterized by mass spectrometry.

Finally, the N-methylation of the peptidyl resins 19–21 was achieved following the three-step procedure described for the preparation of resins 16–18 (Scheme 3). Thus, oNBS protection, Mitsunobu N-methylation, and oNBS group removal afforded the tailored biaryl cyclic peptidyl resins 22–24. Acidolytic cleavage of an aliquot of these resins and HPLC analysis revealed the formation of the expected biaryl cyclic peptides 25–27 in 84, 65, and 32% purity, respectively. Peptides 25–27 were isolated by reverse-phase column chromatography in >99, 78, and 50% HPLC purity, respectively. These compounds were characterized by mass spectrometry and peptide 25 also by NMR spectroscopy.
(HRMS) of these compounds showed a strong peak corresponding to [M + H]⁺. Compounds 31−33 were purified by column chromatography and obtained in 87−97% HPLC purity. Biaryl cyclic cores 31 and 32 were also fully characterized by NMR spectroscopy. These results demonstrated that the presence of an amino acid at the N-methylated tail of these compounds facilitates the incorporation of the palmitic acid.

**CONCLUSIONS**

The first total solid-phase synthesis of arylomycin derivatives incorporating a Phe−Tyr, a Tyr−Tyr, or a His−Tyr biaryl linkage is reported. The biaryl cyclic cores were synthesized on solid support through Miyaura borylation of a 3-iodotyrosine-containing peptidyl resin, subsequent coupling of the corresponding haloamino acid, and final cyclization via an intramolecular Suzuki−Miyaura cross-coupling reaction. The lipopeptidyl tail was assembled through N-methylation, coupling of the corresponding amino acids, and acylation. The characterization of the arylomycin derivatives containing an extra lysine at the lipopeptidyl tail confirmed the effectiveness of our strategy. In this synthetic approach, all steps are performed on the solid phase, including the preparation of the boronotyrosine and the Suzuki−Miyaura reaction. Thus, it not only offers the advantages of the solid phase but can also allow the synthesis of other biaryl-containing natural products as well as the preparation of collections of synthetic arylomycin analogues and, therefore, the identification of new bioactive compounds.

**EXPERIMENTAL SECTION**

**General Methods.** Manual peptide synthesis was performed in polypropylene syringes (2 or 5 mL) fitted with a polyethylene porous disk. Solvents and soluble reagents were removed by suction. Most chemicals were purchased from commercial suppliers Merck (Madrid, Spain), Iris Biotech GmbH (Marktredwitz, Germany), Scharlab (Sentmenat, Spain), or Panreac (Castellar del Vallés, Spain) and used without further purification.

Peptides were analyzed under standard analytical HPLC conditions with a Dionex liquid chromatography instrument composed of a UV/Vis Dionex UVD170U detector, a P680 Dionex bomb, an ASI-100 Dionex automatic injector, and CHROMELION 6.60 software. Detection was performed at a wavelength of 220 nm. Solvent A was 0.1% aqueous TFA, and solvent B was 0.1% TFA in CH₃CN. Conditions A: Analysis was carried out with a 150 mm × 4.6 mm, 5 μm column with a 2−100% B over 7 min at a flow rate of 1 mL/min. Conditions B: Analysis was carried out with a 150 mm × 4.6 mm, 5 μm column with a 2−100% B over 28 min at a flow rate of 1 mL/min.

All purifications were performed on a CombiFlash R200 automated flash chromatography system using RediSep RF Gold reverse-phase C₁₈ column packed with high-performance C₁₈ derivatized silica.

ESI-MS analyses were performed at the Serveis Tècnics de Recerca of the University of Girona with an Esquire 6000 ESI ion trap LC/MS (Bruker Daltonics) instrument equipped with an electrospray ion source. The instrument was operated in the positive ESI(+) ion mode. Samples (5 μL) were introduced into the mass spectrometer ion source directly through an HPLC autosampler. The mobile phase (80:20 CH₃CN/H₂O at a flow rate of 100 μL/min) was delivered by a 1200 Series HPLC pump (Agilent). Nitrogen was employed as both the drying and nebulizing gas.

HRMS was performed on a Bruker MicroTof-QITM instrument using an ESI ionization source at the Serveis Tècnics de Recerca of the University of Girona. Samples were introduced into the mass spectrometer ion source by direct infusion using a syringe pump and were externally calibrated using sodium formate. The instrument was operated in the positive ion mode.

H and ¹³C NMR spectra were recorded with a Bruker 300 or 400 MHz NMR spectrometer at the Serveis Tècnics de Recerca of the University of Girona. Chemical shifts were reported as δ values (ppm) directly referenced to the solvent signal.

Microwave-assisted reactions were performed with a single-mode Discover S-Class lab station microwave (CEM) (0−300 W). The time, temperature, and power were controlled with the Synergy software. The temperature was monitored through an infrared sensor in the floor of the cavity.

**Synthesis of Tr-Ala-Tyr(3-I,Me)-Rink-MBHA (11).** Fmoc-Rink-MBHA resin (0.4 mmol/g) was placed in a polypropylene syringe fitted with a polyethylene filter disk and swollen with CH₂Cl₂ (1 × 20 min) and DMF (1 × 20 min), treated with piperidine/DMF (3:7, 1 × 5 min), and washed with DMF (6 × 1 min). Then, removal of the Fmoc protecting group was achieved with a mixture of piperidine/DMF (3:7, 2 × 10 min), followed by washes with DMF (6 × 1 min). Fmoc-Tyr(3-I,Me)-OH⁵⁶ (2 equiv) was coupled in the presence of COMU (2 equiv), Oxyma (2 equiv), and DIEA (4 equiv) in DMF under stirring at room temperature overnight. The completion of the reaction was monitored by the Kaiser test.⁴⁴ The resin was then washed with DMF (6 × 1 min) and CH₂Cl₂ (3 × 1 min). After Fmoc removal as described above, coupling of commercially available Fmoc-Ala-OH (4 equiv) was performed using DIPCDI (4 equiv) and Oxyma (4 equiv) in DMF at room temperature for 1 h, followed by washes with DMF (6 × 1 min) and CH₂Cl₂ (3 × 1 min). The completion of the reaction was monitored by the Kaiser test.⁴⁴ An aliquot of the resulting Fmoc-protected iodopeptidyl resin 10 was cleaved with TFA/H₂O/TIS (95:2.5:2.5) while stirring at room temperature for 2 h. Following TFA evaporation and diethyl ether extraction, the crude peptide was dissolved in H₂O/CH₃CN (1:1) and lyophillized to give Fmoc-Ala-Tyr(3-I,Me)-NH₂ in 98% purity. τ₀ = 8.51 min (Conditions A).

Next, the N-terminal Fmoc group of 10 was removed as described above, and a trytyl group was introduced using TrCl (10 equiv) and DIEA (10 equiv) in DMF under stirring at room temperature for 4 h. Then, the resin was washed with DMF (6 × 1 min), CH₂Cl₂ (3 × 1 min), and diethyl ether (3 × 1 min). The completion of this reaction was monitored with the Kaiser test.⁴⁴ Acidolytic cleavage of an aliquot of the resulting iodopeptidyl resin 11 afforded H-Ala-Tyr(3-I,Me)-NH₂ in 92% purity. τ₀ = 6.60 min (Conditions A). MS (ESI): [M + Na]⁺ = 394.1 [M + Na]⁺, 414.0 [M + Na]⁺.

**Synthesis of Tr-Ala-Tyr(3-Bpin,Me)-Rink-MBHA (12).** A 5–10 mL round-bottomed flask was charged with the iodopeptidyl resin 11, bis(pinacolato)boron (B(Pin)₂) (4 equiv), PdCl₂(dppf) (0.18 equiv), and 1,1′-bis-(diphenylphosphanyl)ferrocene (dppf) (0.09 equiv). A thoroughly sonicated solution of KOAc in degassed anhydrous DMSO (20 μL/mg of resin) was then added, and the mixture was heated at 80 °C for 8 h. Upon completion of
the reaction, the resin was washed with DMSO (6 × 1 min), MeOH (6 × 1 min), CH₂Cl₂ (6 × 1 min), and diethyl ether (3 × 1 min). An aliquot of the resulting boronopeptidyl resin 12 was cleaved with TFA/H₂O/TIS (95:2.5:2.5) while stirring for 2 h at room temperature. Following TFA evaporation and diethyl ether extraction, the crude peptide was dissolved in H₂O/CH₂CN (1:1), lyophilized, analyzed by HPLC, and characterized by mass spectrometry. H-Ala-Tyr(3-B(OH)₂,Me)-NH₂ was obtained in 72% purity, which resulted from the hydrolysis of the pinacol boronic ester during HPLC analysis. tᵣ = 13.79 min (Conditions B). MS (ESI): m/z = 310.1 [M + H]⁺, 332.1 [M + Na]⁺.

**General Procedure for the Synthesis of Tripeptidyl Resins 7–9.** The boronopeptidyl resin 12 was swelled with CH₂Cl₂ (1 × 20 min); treated with TFA/H₂O/CH₂Cl₂ (0.2:1:98.8, 2 × 1 min + 3 × 20 min); and washed with DMSO (3 × 1 min), DIEA/CH₂Cl₂ (1:19, 3 × 1 min), CH₂Cl₂ (3 × 1 min), and DMF (3 × 1 min). Then, coupling of Boc-Phe(4-I)-OH with Boc-Tyr(3-B(OH)₂,Me)-NH₂ was obtained in 72% purity, which resulted from the hydrolysis of the pinacol boronic ester during HPLC analysis. tᵣ = 13.79 min (Conditions B). MS (ESI): m/z = 525.1, 527.1 [M + H]⁺, 547.1, 549.1 [M + Na]⁺.

**Synthesis of Boc-Phe(4-I)-Ala-Tyr(3-BPin,Me)-Rink-MBHA (7).** This peptidyl resin was prepared using the procedure described above, employing Boc-Phe(4-I)-OH.37 Acidolytic cleavage of the resulting resin 7 for 2 h afforded H-Phe(4-I)-Ala-Tyr(3-B(OH)₂,Me)-NH₂ in 81% purity, resulting from the hydrolysis of the pinacol boronic ester during HPLC analysis. tᵣ = 18.03 min (Conditions B). MS (ESI): m/z = 583.1 [M + H]⁺.

**Synthesis of Boc-Phe(4-I)-Ala-Tyr(3-BPin,Me)-Rink-MBHA (8).** This peptidyl resin was prepared using the procedure described above, employing Boc-Phe(4-I)-OH.37 Acidolytic cleavage of the resulting resin 8 for 2 h afforded H-Tyr(3-I,Me)-Ala-Tyr(3-B(OH)₂,Me)-NH₂ in 82% purity, resulting from the hydrolysis of the pinacol boronic ester during HPLC analysis. tᵣ = 17.99 min (Conditions B). MS (ESI): m/z = 613.1 [M + H]⁺, 635.1 [M + Na]⁺.

**Synthesis of Boc-His(5-Br,1-SEM)-Ala-Tyr(3-BPin,Me)-Rink-MBHA (9a) and Boc-His(5-Br,3-SEM)-Ala-Tyr(3-BPin,Me)-Rink-MBHA (9b).** These peptidyl resins were prepared using the procedure described above, employing a regiosomeric mixture of Boc-His(5-Br,1-SEM)-OH and Boc-His(5-Br,3-SEM)-OH.30 Acidolytic cleavage of the resulting resins 9 for 3 h afforded H-His(5-Br)-Ala-Tyr(3-B(OH)₂,Me)-NH₂ in 82% purity, resulting from the hydrolysis of the pinacol boronic ester during HPLC analysis. tᵣ = 14.20 min (Conditions B). MS (ESI): m/z = 525.1, 527.1 [M + H]⁺, 547.1, 549.1 [M + Na]⁺.

**Synthesis of the Biaryl Cyclic Tripeptidyl Resins 4–6.**

**General Method for the Microwave-Assisted Solid-Phase Intramolecular Suzuki–Miura Arylation.** A 15 mL reaction vessel containing a magnetic stir bar was charged with the corresponding linear peptidyl resin 7–9 (100 mg), Pd₂(dba)₃ (0.2 equiv), SPhos (0.4 equiv), and KF (4 equiv). Thoroughly degassed DME/MeOH/H₂O (9:9:2, 0.8 mL) was then added under nitrogen. The reaction mixture was heated using microwave irradiation at 120 or 140 °C for 30 min. Then, upon cooling, the solvent was removed by filtration and the resin was washed with DMF (6 × 1 min), EtOH (6 × 1 min), CH₂Cl₂ (6 × 1 min), and diethyl ether (3 × 1 min). A solution of TFA/H₂O/TIS (95:2.5:2.5) was then added to an aliquot of the resulting resin, and the mixture was stirred at room temperature for 2 or 3 h. TFA evaporation and diethyl ether extraction gave the corresponding crude peptide mixture, which was dissolved in H₂O/CH₂CN (1:1), lyophilized, analyzed by HPLC, characterized by mass spectrometry, and purified by reverse-phase column chromatography. Pure biaryl cyclic tripeptides were analyzed by HPLC, and characterized by mass spectrometry and NMR spectroscopy.

**Biaryl Cyclic Tripeptidyl Resin 4.** This biaryl cyclic tripeptidyl resin was prepared from the linear peptidyl resin Boc-Phe(4-I)-Ala-Tyr(3-BPin,Me)-Rink-MBHA (7) following the procedure described above at 120 °C. Acidolytic cleavage of an aliquot of the resulting resin 4 for 2 h afforded the biaryl cyclic tripeptide 13 in 92% purity. tᵣ = 16.75 min (Conditions B). MS (ESI): m/z = 411.1 [M + H]⁺. Reverse-phase column chromatography and elution with H₂O/CH₂CN (80:20) afforded the expected biaryl cyclic tripeptide 13 in 99% purity. tᵣ = 6.20 min (Conditions A).1H NMR (400 MHz, CDOD): δ = 7.41 [dd, J = 1.6 and 8.0 Hz, 1H, CH₅aronPhe], 7.34 [dd, J = 1.6 and 8.0 Hz, 1H, CH₅aronPhe], 7.22 [dd, J = 1.6 and 8.0 Hz, 1H, CH₅aronPhe], 7.12–7.08 [m, 2H, CH₂aronPhe], 6.97 [d, J = 8.4 Hz, 1H, CH₅aronTyr], 6.81 [d, J = 2.4 Hz, 1H, CH-2aronTyr], 4.31 [dd, J = 4.0 and 9.2 Hz, 1H, CH-α], 4.01 [q, J = 6.6 Hz, 1H, CH-αAla], 3.90–3.86 [m, 1H, CH-α], 3.86 [s, 3H, OCH₃], 3.52–3.42 [m, 1H, CH-β], 2.92–2.82 [m, 3H, CH-β], 1.26 [d, J = 6.6 Hz, 3H, CH₃-Ala] ppm.

**Biaryl Cyclic Tripeptidyl Resin 5.** This biaryl cyclic tripeptidyl resin was prepared from the linear peptidyl resin Boc-Tyr(3-I,Me)-Ala-Tyr(3-BPin,Me)-Rink-MBHA (8) following the procedure described above at 120 °C. Acidolytic cleavage of an aliquot of the resulting resin 5 for 2 h afforded the biaryl cyclic tripeptide 14 in 73% purity. tᵣ = 15.97 min (Conditions B). MS (ESI): m/z = 441.2 [M + H]⁺. Reverse-phase column chromatography and elution with H₂O/CH₂CN (85:15) afforded the expected biaryl cyclic tripeptide 14 in >99% purity. tᵣ = 5.98 min (Conditions A).1H NMR (400 MHz, CDOD): δ = 7.21 [dd, J = 2.2 and 8.4 Hz, 1H, CH₅aronTyr], 7.14 [dd, J = 2.2 and 8.4 Hz, 1H, CH₅aronTyr], 7.05 [d, J = 2.2 Hz, 1H, CH-2aronTyr], 6.96 [d, J = 8.4 Hz, 1H, CH-5aronTyr], 6.90 [d, J = 8.4 Hz, 1H, CH-5aronTyr], 6.84 [d, J = 2.2 Hz, 1H, CH-2aronTyr], 6.68 [d, J = 2.0 Hz, 1H, CH-α], 6.62 [d, J = 2.0 Hz, 1H, NH], 4.80–4.75 [m, 1H, CH-αAla], 4.63–4.45 [m, 1H, CH-β], 4.22 [dd, J = 3.6 and 4.0 Hz, 1H, CH-α], 3.77 [s, 3H, OCH₃], 3.72 [s, 3H, OCH₃], 3.39 [dd, J = 3.6 and 15.0 Hz, 1H, CH-β], 3.05 [dd, J = 4.0 and 15.0 Hz, 1H, CH-β], 2.97–2.83 [m, 2H, CH-β], 1.39 [d, J = 6.8 Hz, 3H, CH₃-Ala] ppm. MS (ESI): m/z = 441.2 [M + H]⁺, 463.1 [M + Na]⁺.

**Biaryl Cyclic Tripeptidyl Resin 6.** These biaryl cyclic tripeptidyl resins were prepared from the regiosomeric linear peptidyl resins Boc-His(5-Br,1-SEM)-Ala-Tyr(3-BPin,Me)-Rink-MBHA (9a) and Boc-His(5-Br,3-SEM)-Ala-Tyr(3-
The page contains a detailed chemical synthesis process involving the preparation of a biaryl cyclic tripeptide. The process involves several steps including the preparation of resin, coupling of amino acids, protection and deprotection of amino acids, and finally the cleavage of the desired tripeptide. The synthesis involves the use of Fmoc chemistry for amino acid coupling, protection with Boc and NBS, and deprotection with TFA. The final product was analyzed by HPLC and mass spectrometry. The synthetic procedure is described in detail with specific conditions and yields provided.

**Synthesis of the N-Methylated Biaryl Cyclic Tripeptidyl Resins 16–18. General Procedure for the N-Methylation under Mitsunobu Conditions.** The N-terminal Boc group of the corresponding biaryl cyclic tripeptide resin 4–6 was removed by treatment with a solution of TMSOTf and 2,6-lutidine in CH2Cl2 (final concentrations: 2 M TMSOTf and 3 M 2,6-lutidine) (10 × 30 min) at room temperature, and the resulting resin was washed with CH2Cl2 (5 × 1 min), MeOH (3 × 5 min), and DMF (5 × 1 min). A solution of nBS-Cl (4 equiv) and DIEA (10 equiv) in CH2Cl2 was then added to the resin, and the mixture was shaken for 90 min at room temperature. The resulting nBS-protected peptidyl resin was washed with DMF (3 × 0.5 min), CH2Cl2 (3 × 0.5 min), and diethyl ether (3 × 1 min). An aliquot of this peptidyl resin was cleaved with TFA/H2O/TIS (95:2.5:2.5) for 2 or 3 h, and the crude was analyzed by HPLC and mass spectrometry.

Next, the nBS-protected resin was N-methylated by treatment with a solution of PPh3 (25 equiv) and MeOH (50 equiv) in dry THF, followed by the addition of DIAD (25 equiv) in dry THF. The reaction mixture was stirred at room temperature for 1 h. Then, the solution was filtered, and the resulting resin was washed with THF (3 × 0.5 min), CH2Cl2 (3 × 0.5 min), DMF (3 × 0.5 min), and diethyl ether (3 × 1 min). The resulting resin was cleaved for 2 or 3 h, analyzed by HPLC, and characterized by mass spectrometry.

Finally, the nBS group was selectively removed by treatment with β-mercaptoethanol (10 equiv) and DBU (5 equiv) in DMF (2 × 15 min) at room temperature under stirring, followed by washes with DMF (3 × 0.5 min), CH2Cl2 (3 × 0.5 min), DMF (3 × 0.5 min), CH2Cl2 (3 × 0.5 min), and diethyl ether (3 × 1 min). Acidolytic cleavage of an aliquot of the resulting N-methylated resin for 2 or 3 h afforded the expected N-methylated biaryl cyclic tripeptide, which was analyzed by HPLC and characterized by mass spectrometry.

**N-Methylated Biaryl Cyclic Tripeptidyl Resin 16.** N-Methylated cyclic peptidyl resin 16 was prepared from biaryl cyclic tripeptide resin 4 following the general procedure described above. nBS protection: acidolytic cleavage for 2 h of an aliquot of the resulting resin yielded the expected peptide in 69% purity. tR = 7.21 min (Conditions A). Mitsunobu N-methylation: acidolytic cleavage for 2 h of an aliquot of the resulting resin yielded the expected peptide in 89% purity. tR = 7.84 min (Conditions A). MS (ESI): m/z = 640.4 [M + H]+, 662.4 [M + Na]+.

**Tailed Biaryl Cyclic Hexapeptidyl Resin 19.** This peptidyl resin was prepared from the N-methylated peptidyl resin 16 following the general method described above. Acidolytic cleavage of an aliquot of the resulting resin 19 for 2 h yielded the expected biaryl peptide in 85% purity. tR = 6.32 min (Conditions A). MS (ESI): m/z = 862.9 [M + H]+, 884.9 [M + Na]+.

**Tailed Biaryl Cyclic Hexapeptidyl Resin 20.** This peptidyl resin was prepared from the N-methylated peptidyl resin 17 following the general method described above. Acidolytic cleavage of an aliquot of the resulting resin 20 for 2 h yielded the expected biaryl peptide in 85% purity. tR = 6.32 min (Conditions A). MS (ESI): m/z = 862.9 [M + H]+, 884.9 [M + Na]+.
cleavage of an aliquot of the resulting resin 20 for 2 h yielded the expected biaryl peptide in 75% purity. *t*<sub>R</sub> = 6.07 min (Conditions A). MS (ESI): *m/z* = 670.3 [M + H]<sup>+</sup>, 692.2 [M + Na]<sup>+</sup>.

**Regioisomeric Tailed Biaryl Cyclic Hexapeptidyl Resins 21.** These regioisomeric peptidyl resins were prepared from the regioisomeric N-methylated peptidyl resins 18 following the general method described above. Acidolytic cleavage of an aliquot of the resulting resins 21 for 3 h yielded the expected biaryl peptide in 18% purity. *t*<sub>R</sub> = 5.03 min (Conditions A). MS (ESI): *m/z* = 630.3 [M + H]<sup>+</sup>, 652.3 [M + Na]<sup>+</sup>.

**Synthesis of the N-Methylated Tailed Biaryl Cyclic Hexapeptidyl Resins 22–24.** N-Methylation of peptidyl resins 19–21 was performed following the general procedure described for 16–18. After cleavage of an aliquot of the resulting N-methylated resin for 2 or 3 h, the crude peptide was analyzed by HPLC, characterized by mass spectrometry, and purified by reverse-phase column chromatography. Pure tailed biaryl cyclic peptides were analyzed by HPLC, and characterized by HRMS.

**N-Methylated Tailed Biaryl Cyclic Hexapeptidyl Resin 22.** N-methylated peptidyl resin 22 was prepared from the peptidyl resin 19 following the general procedure described for 16–18. oNBS protection: acidolytic cleavage for 2 h of an aliquot of the resulting resin afforded the expected peptide in 90% purity. *t*<sub>R</sub> = 7.10 min (Conditions A). MS (ESI): *m/z* = 825.3 [M + H]<sup>+</sup>, 847.3 [M + Na]<sup>+</sup>. Mitsunobu N-methylation: acidolytic cleavage for 2 h of an aliquot of the resulting resin afforded the expected peptide in 87% purity. *t*<sub>R</sub> = 7.26 min (Conditions A). MS (ESI): *m/z* = 839.3 [M + H]<sup>+</sup>, 861.3 [M + Na]<sup>+</sup>. oNBS group removal: acidolytic cleavage for 2 h of an aliquot of the resulting resin 22 afforded the expected N-methylated tailed biaryl cyclic peptide 25 in 84% purity. *t*<sub>R</sub> = 6.39 min (Conditions A). MS (ESI): *m/z* = 654.3 [M + H]<sup>+</sup>, 676.3 [M + Na]<sup>+</sup>. Reverse-phase column chromatography and elution with H<sub>2</sub>O/CH<sub>3</sub>CN (70:30) afforded the expected biaryl cyclic peptide 25 in >99% purity. *t*<sub>R</sub> = 6.45 min (Conditions A). <sup>1</sup>H NMR (400 MHz, CD<sub>3</sub>OD): δ = 3.78 [dd, *J* = 1.6 and 8.0 Hz, 1H, CH<sub>α</sub>-εamideTyr], 7.29 [dd, *J* = 1.6 and 8.0 Hz, 1H, CH<sub>ε</sub>-amideTyr], 7.25 [dd, *J* = 1.6 and 8.0 Hz, 1H, CH<sub>ε</sub>-宗Phe], 7.15 [dd, *J* = 1.6 and 8.0 Hz, 1H, CH<sub>α</sub>-εamideTyr], 7.20 [dd, *J* = 2.2 and 8.4 Hz, 1H, CH<sub>ε</sub>-宗Phe], 6.96 [dd, *J* = 1.6 and 8.0 Hz, 1H, CH<sub>ε</sub>-宗Phe], 6.79 [dd, *J* = 2.2 and 8.4 Hz, 1H, CH<sub>ε</sub>-宗Phe], 6.65 [dd, *J* = 1.6 and 8.0 Hz, 1H, CH<sub>ε</sub>-宗Phe], 6.79 [dd, *J* = 8.4 Hz, 1H, CH<sub>α</sub>-宗Phe], 6.23 [dd, *J* = 8.4 Hz, 1H, CH<sub>ε</sub>-宗Phe], 6.07 [dd, *J* = 1.6 and 8.0 Hz, 1H, CH<sub>ε</sub>-宗Phe]. MS (ESI): *m/z* = 444.3 [M + H]<sup>+</sup>, 466.2 [M + Na]<sup>+</sup>. Reverse-phase column chromatography and elution with H<sub>2</sub>O/CH<sub>3</sub>CN (98:2) afforded the expected biaryl cyclic peptide 27 in 50% purity. *t*<sub>R</sub> = 5.30 min (Conditions A). MS (ESI): *m/z* = 644.3 [M + H]<sup>+</sup>, 666.3 [M + Na]<sup>+</sup>. HRMS (ESI): calcd for C<sub>48</sub>H<sub>74</sub>N<sub>7</sub>O<sub>9</sub> [M + Na]<sup>+</sup> 666.2970, found 666.2954.

**Synthesis of Tailed Biaryl Cyclic Lipohexapeptides 1–3.** **General Method for the Solid-Phase Peptide Acylation of N-Methylated Tailed Biaryl Cyclic Hexapeptidyl Resins.** A solution of palmitic acid (4 equiv), DIPCDI (4 equiv), and Oxyma (4 equiv) in DMSO was added to the corresponding N-methylated tailed biaryl cyclic hexapeptidyl resin 22–24. The reaction mixture was stirred overnight at room temperature. After acylation, the reaction resin was washed with DMSO (3 × 1 min), CH<sub>2</sub>Cl<sub>2</sub> (3 × 1 min), and diethyl ether (3 × 1 min), and the completion of the reaction was then checked using the chloranil test. Upon completion of the acylation, a solution of TFA/H<sub>2</sub>O/TIS (95:2.5:2.5) was added to the resin and the completion of the reaction was then checked using the chloranil test. Upon completion of the acylation, a solution of TFA/H<sub>2</sub>O/TIS (95:2.5:2.5) was added to the resin and the completion of the reaction was then checked using the chloranil test. Upon completion of the reaction, the reaction mixture was stirred overnight at room temperature. After acylation, the reaction resin was washed with DMSO (3 × 1 min), CH<sub>2</sub>Cl<sub>2</sub> (3 × 1 min), and diethyl ether (3 × 1 min), and the completion of the reaction was then checked using the chloranil test. Upon completion of the reaction, the reaction mixture was stirred overnight at room temperature. After acylation, the reaction resin was washed with DMSO (3 × 1 min), CH<sub>2</sub>Cl<sub>2</sub> (3 × 1 min), and diethyl ether (3 × 1 min), and the completion of the reaction was then checked using the chloranil test. Upon completion of the reaction, the reaction mixture was stirred overnight at room temperature. After acylation, the reaction resin was washed with DMSO (3 × 1 min), CH<sub>2</sub>Cl<sub>2</sub> (3 × 1 min), and diethyl ether (3 × 1 min), and the completion of the reaction was then checked using the chloranil test. Upon completion of the reaction, the reaction mixture was stirred overnight at room temperature. After acylation, the reaction resin was washed with DMSO (3 × 1 min), CH<sub>2</sub>Cl<sub>2</sub> (3 × 1 min), and diethyl ether (3 × 1 min), and the completion of the reaction was then checked using the chloranil test.

**Tailed Biaryl Cyclic Lipohexapeptide 1.** This peptide was prepared from resin 22 following the method described above. Acidolytic cleavage of an aliquot of the resulting lipopeptidyl resin for 2 h gave traces of the expected tailed biaryl cyclic lipopeptide 1. HRMS (ESI): calcd for C<sub>60</sub>H<sub>86</sub>N<sub>12</sub>O<sub>10</sub> [M + H]<sup>+</sup> 892.5543, found 892.5549.

**Tailed Biaryl Cyclic Lipohexapeptide 2.** This peptide was prepared from resin 23 following the method described above. Acidolytic cleavage of an aliquot of the resulting lipopeptidyl resin for 2 h gave traces of the expected tailed biaryl cyclic lipopeptide 2. MS (ESI): *m/z* = 922.6 [M + H]<sup>+</sup>. This peptide was prepared from resin 24 following the method described above.
Acidolytic cleavage of an aliquot of the resulting lipopeptidyl cyclic peptide 3 in 3 h gave traces of the expected tailed biaryl cyclic lipopeptide. HRMS (ESI): calcd for C\textsubscript{45}H\textsubscript{72}N\textsubscript{9}O\textsubscript{9} [M + H]\textsuperscript{+} 702.1286, found 702.1275; calcd for C\textsubscript{51}H\textsubscript{84}N\textsubscript{11}O\textsubscript{10} [M + Na]\textsuperscript{+} 724.5251, found 724.5248.

### 2.24 \[t, \text{CH}_3\text{CN} (55:45)\] yielded pure calcd for C\textsubscript{54}H\textsubscript{86}N\textsubscript{9}O\textsubscript{10} [M + H]\textsuperscript{+} 1020.6492, found 1020.6484; calcd for C\textsubscript{54}H\textsubscript{85}N\textsubscript{9}O\textsubscript{10}Na [M + Na]\textsuperscript{+} 1042.6312, found 1042.6281.

#### **Synthesis of Tailed Biaryl Cyclic Lipopeptides**

**31–33. General Method for the Solid-Phase Peptide Synthesis of Tailed Biaryl Cyclic Lipopeptide Resins.** A solution of Fmoc-Lys(Boc)-OH (4 equiv), DIPCDI (4 equiv), and Oxya (4 equiv) in DMF was added to the corresponding N-methylated resins and stirred at room temperature for 2 or 3 h. Following TFA evaporation and diethyl ether extraction, the crude peptide was dissolved in H\textsubscript{2}O/CH\textsubscript{3}CN (1:1), lyophilized, and characterized by mass spectrometry. Tailed biaryl cyclic lipopeptides were purified by reverse-phase column chromatography.

**Tailed Biaryl Cyclic Lipopeptide 31.** This biaryl peptide was prepared from peptidyl resin 22 as described above. After coupling of Fmoc-Lys(Boc)-OH, acidolytic cleavage of an aliquot of the resulting resin 28 for 2 h afforded the expected biaryl peptide in 81% purity. t\textsubscript{R} = 7.36 min (Conditions A). MS (ESI): m/z = 1004.6 [M + H]\textsuperscript{+}, 1026.5 [M + Na]\textsuperscript{+}. Coupling of palmitic acid and acidolytic cleavage for 2 h afforded the expected biaryl cyclic palmitoyl peptide 31 in 69% purity. t\textsubscript{R} = 9.13 min (Conditions A). MS (ESI): m/z = 1020.7 [M + H]\textsuperscript{+}, 1042.7 [M + Na]\textsuperscript{+}. Elution with H\textsubscript{2}O/CH\textsubscript{3}CN (55:45) yielded pure 31 in 97% purity. t\textsubscript{R} = 9.18 min (Conditions A). MS (ESI): m/z = 1010.8 [M + H]\textsuperscript{+}, 1032.7 [M + Na]\textsuperscript{+}. HRMS (ESI): calcd for C\textsubscript{51}H\textsubscript{85}N\textsubscript{11}O\textsubscript{10} [M + H]\textsuperscript{+} 1050.7 [M + Na]\textsuperscript{+}. HRMS (ESI): calcd for C\textsubscript{51}H\textsubscript{85}N\textsubscript{11}O\textsubscript{10}Na [M + Na]\textsuperscript{+} 1072.7 [M + Na]\textsuperscript{+}. Elution with H\textsubscript{2}O/CH\textsubscript{3}CN (55:45) yielded pure 33 in 87% purity. t\textsubscript{R} = 8.67 min (Conditions A). MS (ESI): m/z = 505.8 [M + 2H]\textsuperscript{2+}, 1010.7 [M + H]\textsuperscript{+}, 1032.7 [M + Na]\textsuperscript{+}.

**Supporting Information**

The Supporting Information is available free of charge at https://pubs.acs.org/doi/10.1021/acsomega.0c03352.

HPLC and mass spectrometry of linear peptides, biaryl cyclic peptides, and biaryl cyclic lipopeptides. 1D and 2D NMR of biaryl cyclic tripeptides 13–15, N-methylated tailed biaryl cyclic hexapeptide 25, and tailed biaryl cyclic lipopeptides 31 and 32 (PDF)

#### **ASSOCIATED CONTENT**

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