

# Total Synthesis of the Highly *N*-Methylated Acetylene-Containing Anticancer Peptide Jahanyne

Andrew Siow,<sup>†</sup> George Opiyo,<sup>‡</sup> Iman Kavianinia,<sup>†,§</sup> Freda F. Li,<sup>‡</sup> Daniel P. Furkert,<sup>§,‡</sup> Paul W. R. Harris,<sup>†,§</sup> and Margaret A. Brimble<sup>\*,†,§,‡</sup>

<sup>†</sup>School of Biological Sciences, The University of Auckland, 3A Symonds Street, Auckland 1010, New Zealand <sup>‡</sup>School of Chemical Sciences, The University of Auckland, 23 Symonds Street, Auckland 1010, New Zealand <sup>§</sup>Maurice Wilkins Centre for Molecular Biodiscovery, The University of Auckland, 3A Symonds Street, Auckland 1010, New Zealand

**(5)** Supporting Information

**ABSTRACT:** The first total synthesis of the highly *N*-methylated acetylene-containing lipopeptide jahanyne, an apoptosis-inducing natural product from marine cyanobacteria, is reported. A late-stage solution-phase coupling enabled introduction of the *C*-terminal ketone pyrrolidine moiety. A modified Fmoc solid-phase synthesis strategy was adopted to effectively couple multiple sterically hindered *N*-methylated amino acids while suppressing epimerization. The total synthesis has enabled confirmation of the proposed absolute configuration of natural jahanyne.

arine cyanobacteria are a well-known source of structurally unique bioactive molecules.<sup>1,2</sup> The biological activity of these intriguing natural products ranges from anticancer to antiviral activity, which makes these secondary metabolites attractive leads for biomedical research and drug discovery.<sup>2</sup> Several of these compounds have been isolated, such as dolastatin-10. Synthetic analogues of dolastatin-10 are clinically used as payloads in antibody-drug conjugates for cancer therapy.<sup>3</sup> A small subset of these secondary metabolites share an unusual terminal alkynyl fatty acid motif, including dragonamide, kurahyne, viridamides, carmabin, almiramides, and jahanyne.<sup>4-9</sup> These heavily N-methylated acetylenecontaining lipopeptides exhibit a range of antimicrobial, antifungal, and antitumor activities; hence, further studies enabled by the chemical synthesis of this subfamily of lipopeptides may afford new leads for drug discovery.

In early 2015, Suenaga et al.9 reported the isolation and characterization of a novel acetylene-containing lipopeptide jahanyne (1a) from the marine cyanobacterium Lyngbya sp. (Scheme 1). Jahanyne (1a) contains the rare N-terminal acetylene-containing fatty acid (2R,4S)-2,4-dimethyldec-9-ynoic acid (2), a C-terminal (S)-1-(pyrrolidin-2-yl)ethan-1-one (Oep) (3), and multiple N-methylated amino acids. The substitution of canonical amino acids for N-methylated amino acids has been shown to alter biological functions of polypeptides such as improved stability, binding, and oral bioavailability.<sup>10-12</sup> This makes efficient incorporation of Nmethyl amino acids of paramount importance for peptide-based drug discovery programs. However, the lack of efficient methods to synthesize highly N-methylated and acetylenecontaining lipopeptides hampers detailed investigation of the properties and the biomedical potential of this family of







secondary metabolites. To the best of our knowledge, no member of this family of acetylene-containing *N*-methylated lipopeptides has been synthesized using a 9-fluorenylmethoxy-carbonyl (Fmoc)-solid-phase approach. Currently, the only methodology used to synthesize this class of lipopeptides is a solution-based approach.<sup>4,5</sup>

Received: December 17, 2017

Herein, we describe the first total synthesis of jahanyne (1a) using a combination of solid-phase peptide synthesis and a late stage solution-phase coupling. It was envisaged that the total synthesis of 1a would be achieved by a late stage convergent solution-phase coupling of Oep 3 with the C-terminal fragment 4a (Scheme 1). Conceivably, intermediate 4a would be synthesized by the solid-phase coupling of 5 with the N-terminal building block 2, followed by mild linker cleavage. Assembly of the peptide chain 5 would be accomplished by a linear Fmoc SPPS approach on resin.

We envisaged that the required fatty acid 2 would be prepared in six steps from commercially available 5-hexyn-1-ol (6) (Scheme 2). Hence, the synthesis began by converting





alcohol **6** to iodide 7 via an Appel reaction.<sup>13</sup> Asymmetric alkylation of the enolate generated from **8** with iodide 7, under Myers conditions,<sup>14</sup> proceeded smoothly to afford amide **9**. Reductive cleavage of the chiral auxiliary in **9** afforded alcohol **10** which was then converted to **11**. Iodide **11** was employed in asymmetric alkylation of the enolate generated from *ent-8* to afford **12** thus installing the required 2,4-*anti*-dimethyl stereochemistry in the alkyl chain of amide **12**. Finally, basic hydrolysis of pseudoephedrine amide **12** afforded the target carboxylic acid **2** (dr = *ca*. 15:1). To confirm the absolute stereochemistry, the (*S*)-phenylglycine amide of **2**, prepared in the original isolation paper from **2** obtained by degradation of

Scheme 3. Total Synthesis of Jahanyne (1a) and Analogue 1b

the isolated natural product sample, was also synthesized (see Supporting Information). The spectral data of both samples were in excellent agreement, confirming that the fatty acid moiety of the natural product possessed the 2,4-*anti*, (2*R*,4*S*) configuration.<sup>9</sup>

The requisite C-terminal Oep 3 was synthesized following a literature protocol.  $^{15}$ 

With the necessary building blocks in hand, we then focused on elaboration of the *N*-methylated peptide sequence. The particular challenge of synthesizing *N*-methylated peptides is their poor reactivity in SPPS due to steric hindrance imposed by the additional methyl group.<sup>11,16</sup> Yields are additionally compromised by diketopiperazine (DKP) formation which is prevalent in both the *N*-methyl amino acid coupling and Fmocremoval steps.<sup>16,17</sup> Furthermore, epimerization at the  $\alpha$ -C position of activated *N*-methyl amino acids during the coupling step is also problematic.<sup>11,16</sup>

We avoided using the *tert*-butyloxycarbonyl (Boc) protecting group strategy for SPPS, due to the documented acid lability of *N*-methylated amino acids, which are prone to fragment as well as undergo DKP formation.<sup>17,18</sup> We therefore examined linkers suitable for Fmoc SPPS that can be cleaved under mild conditions and have sufficient steric hindrance on the solid support to help suppress DKP formation.

Our first synthetic approach focused on using the 2chlorotrityl-functionalized polystyrene resin (2-CTC) preloaded with Pro (0.62 mmol/g) at elevated temperatures to promote coupling of hindered amino acids. Initial results indicated that 2-CTC was unstable to microwave (MW) conditions and the high temperature used in MW assisted couplings led to hydrolysis of the chlorotrityl ester bond resulting in premature cleavage of the peptide from the *C*terminus (see pp S16–S19 in Supporting Information).<sup>19</sup>

Given our findings using 2-CTC, we next investigated the use of the mild acid labile 4-(4-hydroxymethyl-3-methoxyphenoxy)butyric acid (HMPB) linker (Scheme 3). We attached Fmoc-Pro-OH to preformed HMPB-aminomethyl polystyrene resin 13 and obtained an acceptable loading of 0.4 mmol/g (66%) based on the Fmoc-release assay (see p S8 in the Supporting Information). The resulting HMPB handle of 14 is cleavable under mild conditions using hexafluoroisopropanol (HFIP) facilitating the ability to monitor the reaction by



DOI: 10.1021/acs.orglett.7b03925 Org. Lett. XXXX, XXX, XXX–XXX HPLC/MS analysis. Removal of the Fmoc group from the Fmoc-N-methylated amino acids was carried out using the hindered Fmoc-deblocking reagent 2-methylpiperidine (2-MP).<sup>20–22</sup> This was used as a precautionary measure to prevent the risk of DKP formation or epimerization of sensitive amino acids.<sup>23,24</sup> Pleasingly, 2-MP quantitatively removed the Fmoc-protecting group, in an acceptable reaction time (1 × 5 min, 1 × 7 min) with no detectable side reactions, and was used for all subsequent Fmoc-removals.

Attention next turned to the optimal choice of coupling reagents to efficiently couple sequential sterically hindered Fmoc-protected N-methylated amino acids. For the majority of the jahanyne sequence, a mixture of (1-cyano-2-ethoxy-2oxoethylidenaminooxy) dimethylaminomorpholinocarbenium hexafluorophosphate (COMU) and ethyl 2-cyano-2-(hydroxyimino)acetate (Oxyma) was used as previously reported to facilitate hindered amide bond formation.<sup>25,26</sup> Epimerization of 15 at the Phe residue from the COMU/ Oxyma mediated coupling between Fmoc-NMe-Phe-OH and resin bound 14 could be negated by using 2,4,6-trimethylpyridine (sym-collidine) as a weaker base (see Figure S7 in the Supporting Information). However, use of MW irradiation was critical to expedite the amidation of 14 to 15 and was used for the following coupling steps involving Fmoc-protected Nmethylated amino acids.<sup>27</sup> Fmoc-removal of 15 resulted in a mixture of HPLC separable peaks with identical mass, which we attributed to cis/trans amide conformers.<sup>17,27,28</sup> This isomerism could be controlled by heating the HPLC column to 77 °C resulting in a single peak due to accelerated cis/trans interconversion, thereby enabling accurate monitoring of reactions (see Figures S8–S9 in the Supporting Information).<sup>27</sup>

Next, we focused on the synthesis of the adjacent sterically demanding *NM*e-Val sequence. Attempts to couple the first Fmoc-*NM*e-Val-OH to deprotected **15** using COMU/Oxyma with *sym*-collidine under MW irradiation resulted in low coupling efficiency. Use of *N*,*N*-diisopropylethylamine (DIPEA) not only quantitatively coupled Fmoc-*NM*e-Val-OH but also promoted epimerization. Employing *N*-methylmorpholine (NMM) and COMU/Oxyma cleanly yielded the desired Fmoc-protected peptide **16** (see Figures S11–S13 in the Supporting Information).

Disappointingly, despite extensive screening of optimal coupling reagents, coupling of the second and third bulky Fmoc-NMe-Val-OH to peptidyl resin 16 was low yielding (see Table S1 in the Supporting Information). Pleasingly, the HNMe-Val coupling steps were solved by generating the Fmocamino acyl chlorides in situ using bis(trichloromethyl)carbonate (BTC) as reported by Falb et al.<sup>29</sup> Fmoc-NMe-Val-OH in dry THF was activated with BTC, and sym-collidine was then added to neutralize the liberated HCl. DIPEA was also added to resin bound 16 to prevent premature acidic cleavage from the resin as well as promote the ensuing reaction. The activated Fmoc-NMe-Val-Cl was added to the peptidyl resin pretreated with DIPEA and stirred at room temperature for 4 h, resulting in quantitative conversion to 17 with no detectable epimerization.<sup>16,29</sup> The remaining amino acid couplings to afford resin bound 5 were effected using COMU/Oxyma and appropriate base under MW irradiation.

Having established a reliable synthetic method to form the linear backbone of jahanyne, we next investigated coupling of the acetylene-containing fatty acid moiety 2 to the *N*-terminus of **5**. For model studies, we utilized commercially available 10-undecynoic acid (10-UDYA) to couple with peptidyl resin **5** 

using COMU/Oxyma and NMM, with a prolonged reaction time of 3 h at room temperature, followed by HFIP cleavage to afford **4b**. We then successfully appended synthetic fatty acid **2** to peptidyl resin **5** using similar coupling conditions with that of **4b**. Treatment of the fully assembled resin-bound peptide with 20 vol % HFIP in CH<sub>2</sub>Cl<sub>2</sub>, and analysis by HRMS, confirmed the presence of the desired peptide **4a** as the major product (*ca.* 66% as judged by peak area integration of RP-HPLC,  $\lambda = 210$  nm,  $t_{\rm R} = 59.6$  min) (Figure 1). Fragmentation



**Figure 1.** RP-HPLC analysis of crude **4a** using a Gemini C-18 column (5  $\mu$ m, 10 mm × 250 mm). HRMS (EI): m/z [M + H]<sup>+</sup> calculated for C<sub>54</sub>H<sub>86</sub>N<sub>7</sub>O<sub>9</sub><sup>+</sup>: 976.6481, observed: 976.6472; [M + Na]<sup>+</sup> calculated for C<sub>54</sub>H<sub>85</sub>N<sub>7</sub>O<sub>9</sub>Na<sup>+</sup>: 998.6301, observed: 998.6317.

of peptide 4a was observed during HRMS, as a consequence of the labile *NMe-Phe-Pro* and *NMe-Val-NMe-Phe* amide bonds (see p S30 in the Supporting Information). The resulting crude peptide 4a was then directly subjected to the final convergent solution-phase installation of the *C*-terminal Oep moiety.

We anticipated poor reactivity for the final solution-phase coupling of Oep 3 to linear peptide 4a/4b, due to the hydrophobic properties of our heavily *N*-methylated peptide fragment and the need to effect formation of a tertiary amide bond. We therefore conducted model studies using Fmoc-Pro-OH to identify suitable coupling conditions. Initial studies using *N*,*N'*-dicyclohexylcarbodiimide (DCC) or *N*-(3-(dimethylamino)propyl)-*N'*-ethylcarbodiimide hydrochloride (EDC·HCl) to effect the coupling of Oep 3 to Fmoc-Pro-OH were disappointing. However, condensation using *N*,*N'*-diisopropylcarbodiimide (DIC) with 6-Cl-HOBt as an additive using DIPEA as a base was successful.

Unfortunately, extension of these model conditions to effect the desired coupling of Oep with 4a/4b were unsuccessful. It was eventually determined that use of a modified cocktail of DIC/6-Cl-HOBt/DIPEA and COMU/Oxyma with DIPEA activation and MW irradiation successfully effected the hindered condensation of 4a/4b with 3 (see Figures S26– S27 in the Supporting Information). Subsequent purification of the crude reaction mixture by RP-HPLC afforded jahanyne (1a, 47%, > 98% purity,  $t_{\rm R}$  = 54.2 min) (Figure 2). Pleasingly, the spectral data for synthetic 1a were observed to correspond well with those originally reported for the natural product 1a (see pp S43–S62 in the Supporting Information).<sup>9</sup>

In summary, we have developed the first synthesis of enantiopure jahanyne (1a) and an analogue 1b containing an achiral alkynoic acid, via Fmoc SPPS. The synthesis described herein not only confirmed the correct structure for the natural



**Figure 2.** RP-HPLC analysis of purified **1a** using an XTerra MS C-18 column (5  $\mu$ m; 4.6 mm × 150 mm). HRMS (EI): m/z [M + Na]<sup>+</sup> calculated for C<sub>60</sub>H<sub>94</sub>N<sub>8</sub>O<sub>9</sub>Na<sup>+</sup>: 1093.7036, observed: 1093.7032.

product jahanyne (1a), but should also be applicable to the synthesis of other acetylene-containing lipopeptides which contain contiguous *N*-methylated amino acids.<sup>4–8</sup> Moreover, the methodology can be applied more generally to the synthesis of *N*-methylated peptides using an SPPS-based approach.

#### ASSOCIATED CONTENT

#### **Supporting Information**

The Supporting Information is available free of charge on the ACS Publications website at DOI: 10.1021/acs.or-glett.7b03925.

Experimental procedures and spectral data (PDF)

#### AUTHOR INFORMATION

## **Corresponding Author**

\*E-mail: m.brimble@auckland.ac.nz.

ORCID <sup>®</sup>

Daniel P. Furkert: 0000-0001-6286-9105 Margaret A. Brimble: 0000-0002-7086-4096

# Notes

The authors declare no competing financial interest.

## ACKNOWLEDGMENTS

We acknowledge financial support from the Maurice Wilkins Centre for Molecular Biodiscovery and The University of Auckland for the award of a Doctoral Scholarship to G.O. The authors would also like to thank Chloë Marcourt (The University of Auckland) for the synthesis of Oep.

#### REFERENCES

(1) Costa, M.; Garcia, M.; Costa-Rodrigues, J.; Costa, M. S.; Ribeiro, M. J.; Fernandes, M. H.; Barros, P.; Barreiro, A.; Vasconcelos, V.; Martins, R. *Mar. Drugs* **2014**, *12*, 98–114.

(2) Mooberry, S. L.; Leal, R. M.; Tinley, T. L.; Luesch, H.; Moore, R. E.; Corbett, T. H. Int. J. Cancer 2003, 104, 512–521.

(3) Polakis, P. Pharmacol. Rev. 2016, 68, 3–19.

(4) Chen, H.; Feng, Y.; Xu, Z.; Ye, T. *Tetrahedron* **2005**, *61*, 11132–11140.

(5) Okamoto, S.; Iwasaki, A.; Ohno, O.; Suenaga, K. J. Nat. Prod. 2015, 78, 2719–2725.

(6) Simmons, T. L.; Engene, N.; Ureña, L. D.; Romero, L. I.; Ortega-Barría, E.; Gerwick, L.; Gerwick, W. H. *J. Nat. Prod.* **2008**, *71*, 1544– 1550. (7) Hooper, G. J.; Orjala, J.; Schatzman, R. C.; Gerwick, W. H. J. Nat. Prod. **1998**, 61, 529–533.

(8) Sanchez, L. M.; Lopez, D.; Vesely, B. A.; Togna, G. D.; Gerwick, W. H.; Kyle, D. E.; Linington, R. G. *J. Med. Chem.* **2010**, *53*, 4187–4197.

(9) Iwasaki, A.; Ohno, O.; Sumimoto, S.; Ogawa, H.; Nguyen, K. A.; Suenaga, K. *Org. Lett.* **2015**, *17*, 652–655.

(10) Chatterjee, J.; Rechenmacher, F.; Kessler, H. Angew. Chem., Int. Ed. 2013, 52, 254–269.

(11) Nabika, R.; Oishi, S.; Misu, R.; Ohno, H.; Fujii, N. *Bioorg. Med. Chem.* **2014**, *22*, 6156–6162.

(12) Biron, E.; Chatterjee, J.; Ovadia, O.; Langenegger, D.; Brueggen, J.; Hoyer, D.; Schmid, H. A.; Jelinek, R.; Gilon, C.; Hoffman, A.; Kessler, H. Angew. Chem., Int. Ed. 2008, 47, 2595–2599.

(13) Silvi, M.; Arceo, E.; Jurberg, I. D.; Cassani, C.; Melchiorre, P. J. Am. Chem. Soc. **2015**, 137, 6120–6123.

(14) Myers, A. G.; Yang, B. H.; Chen, H.; McKinstry, L.; Kopecky, D. J.; Gleason, J. L. J. Am. Chem. Soc. **1997**, 119, 6496–6511.

(15) Kong, C.; Jana, N.; Driver, T. G. Org. Lett. 2013, 15, 824–827.
(16) Thern, B.; Rudolph, J.; Jung, G. Angew. Chem., Int. Ed. 2002, 41, 2307–2309.

(17) Teixidó, M.; Albericio, F.; Giralt, E. J. Pept. Res. 2005, 65, 153–166.

(18) Vaisar, T.; Urban, J. J. Mass Spectrom. 1998, 33, 505-524.

(19) Echalier, C.; Al-Halifa, S.; Kreiter, A.; Enjalbal, C.; Sanchez, P.; Ronga, L.; Puget, K.; Verdié, P.; Amblard, M.; Martinez, J.; Subra, G. *Amino Acids* **2013**, *45*, 1395–1403.

(20) Hachmann, J.; Lebl, M. J. Comb. Chem. 2006, 8, 149-149.

(21) Ali, A. M.; Taylor, S. D. Angew. Chem., Int. Ed. 2009, 48, 2024–2026.

(22) Ali, A. M.; Taylor, S. D. J. Pept. Sci. 2010, 16, 190-199.

(23) Brieke, C.; Cryle, M. J. Org. Lett. 2014, 16, 2454-2457.

(24) Szantai-Kis, M.; Walters, C. R.; Barrett, T. M.; Hoang, E. M.; Petersson, E. J. *Synlett* **2017**, *28*, 1789–1794.

(25) Siow, A.; Hung, K.-y.; Harris, P. W.; Brimble, M. A. Eur. J. Org. Chem. 2017, 2017, 350-354.

(26) Kavianinia, I.; Kunalingam, L.; Harris, P. W.; Cook, G. M.; Brimble, M. A. Org. Lett. **2016**, *18*, 3878–3881.

(27) Rodríguez, H.; Suarez, M.; Albericio, F. J. Pept. Sci. 2010, 16, 136–140.

(28) Schnitzer, T.; Wennemers, H. J. Am. Chem. Soc. 2017, 139, 15356-15362.

(29) Falb, E.; Yechezkel, T.; Salitra, Y.; Gilon, C. J. Pept. Res. 1999, 53, 507-517.