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Total synthesis of halipeptin A, a potent anti-inflammatory cyclodepsipeptide from a marine sponge

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Abstract—Total synthesis of halipeptin A, a potent anti-inflammatory cyclodepsipeptide, was achieved through proline-catalyzed asymmetric α -oxidation, diastereoselective aldol reaction, silver cyanide-mediated esterification, and macrolactamization. © 2005 Elsevier Ltd. All rights reserved.

Halipeptins A (1) and B $(2)^1$ are novel 16-membered cvclodepsipeptides isolated from the marine sponge Haliclona sp. collected in waters off the Vanuatu Islands by Gomez-Paloma and co-workers in 2001. Halipeptin A is known to show strong anti-inflammatory activity in vivo. In 2002, Gomez-Paloma and co-workers. reported isolation of halipeptin C (3) closely related to 1 and 2 from the same sponge, reexamined the original assignments with a novel oxazetidine ring, and corrected the oxazetidine amino acid to thiazoline amino acid in halipeptins A and B (Fig. 1).² Snider reported confirmation of the above revision based on synthesis of the oxazetidine amino acid.³ Halipeptins consist of (S)-alanine and three unique components, the thiazoline-amino acid ((ala)Thz), N-methyl hydroxyisoleucine (N-MeOH-Ile) (or *N*-MeVal for 3), and 3-hydroxy-2,2,4-trimethyl-7-methoxy(or hydroxy for 2 and 3)decanoic acid (HTMMD or HTMHD). The stereostructure at the C3 and C4 of HTMMD and HTMHD remained to be determined except C7, which was confirmed to be S by the Mosher method using HTMHD. In addition to their potent biological activities, their intriguing structures led several groups to initiate efforts directed toward the total synthesis.⁴⁻⁶ The first total synthesis of this unique cyclodepsipeptide, halipeptin A, was accomplished by Ma and co-workers, leading to the structural confirmation of the revised halipeptins.7 A little later, the Nicolaou group also succeeded in synthesis of halipeptin

Keywords: Halipeptin A; Proline-catalyzed asymmetric α-oxidation; Asymmetric aldol reaction; Cyclodepsipeptide; Macrolactamization.







A and the relatives.⁸ These recent reports prompted us to disclose our efforts on the total synthesis of halipeptin A. Our synthesis includes coupling of the ester and amide segments at the HTMMD/(ala)Thz site and final macrocyclization at the *N*-MeOHIle/Ala site. As part of our studies on synthesis of cyclodepsipeptides with biologically interesting activities, we have already demonstrated synthesis of the *N*-MeOHIle derivative.⁶

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Our study began with convergent synthesis of the ester segment including proline-catalyzed α -oxidation,⁹ diastereoselective aldol condensation, and the Julia coupling as the key steps as shown in Scheme 1. Aldol 4 from Kiyooka's chiral oxazaborolidinone-catalyzed asymmetric aldol condensation¹⁰ was converted to acetonide 5 by reduction of 4 and subsequent protection with 2,2-dimethoxypropane and TsOH, which was subjected to deprotection of the benzyl ether and subsequent oxidation of 6 with DMP¹¹ to give aldehyde 7. The required sulfone 11 was prepared from (S)-pentane-1,2-diol (8), which was synthesized from pentanal by introduction of the (S)-hydroxy function using proline-catalyzed asymmetric α -oxidation and subsequent reductive removal of the anilino group. Synthesis of benzyl ether 10 by selective protection was carried out by a sequence of benzylidene acetalization with benzaldehyde dimethylacetal and reductive cleavage of the acetal to a benzyl ether with DIBAL-H. Thianation of 10 with N-phenyl-5-mercaptotetrazole, DEAD, and

triphenylphosphine,¹² and subsequent oxidation with *m*CPBA gave sulfone 11 in excellent yield. The Julia coupling¹³ of 7 and 11 was performed by using KHMDS¹¹ in DME to produce the HTMMD skeleton 12, which was derivatized to HTMMD building block 13 in four steps by simultaneous hydrogenation of the double bond and the benzyl ether, O-methylation with iodomethane and KHMDS, deprotection of the acetonide, and protection of the primary alcohol with TBSCl. In the Julia coupling, the choice of base was critical for success. Use of lithium hexamethyldisilazide and *n*-butyl lithium, common bases for the Julia coupling, in DME failed to afford the coupling product 12. Esterification of the HTMMD 13 with Cbz-(S)-Ala-OH and EDCI/ DMAP¹¹ provided the ester in 68% yield with slight epimerization of a 3:1 ratio at the alanine residue as the Ma group observed,⁷ which was unfortunately inseparable. However, use of Fmoc-(S)-Ala-Cl in the presence of silver cyanide in toluene accomplished racemization-free esterification to afford ester 14 in 14% yield but the



Scheme 1. Synthesis of the ester segment **16**. Reagents and conditions: (a) LiAlH₄, Et₂O, 23 °C, 5.5 h; (b) 2,2-dimethoxypropane, TsOH, CH₂Cl₂, 23 °C, 15 h, 83% (two steps); (c) H₂, Raney-Ni, MeOH, 23 °C, 14 h, 92%; (d) DMP, CH₂Cl₂, 23 °C, 4 h, 94%; (e) PhNO, (*R*)-proline, CHCl₃, 4 °C, 2 h, then NaBH₄, EtOH, 5 °C, 35 min, >99% ee; (f) H₂, Pd–C, MeOH, 23 °C, 17 h, 94% (two steps); (g) PhCH(OMe)₂, TsOH·H₂O, PhMe, 23 °C, 17.5 h, 77%; (h) DIBAL-H, CH₂Cl₂, -17 to -10 °C, 1.5 h, 94%; (i) 1-phenyl-5-mercapto-tetrazole, DEAD, Ph₃P, THF, 23 °C, 9.5 h, 94%; (j) *m*CPBA, CH₂Cl₂, 0-23 °C, 16 h, 93%; (k) KHMDS, DME, **7**, -65 to 23 °C, 16 h, 63%; (l) H₂, Raney-Ni, MeOH, 23 °C, 11.5 h, 63%; (m) KHMDS, THF, then MeI, -78 to 23 °C, 5 h, 86%; (n) 6 M HCl–MeOH (4:1), 0-23 °C, 3 h, 100%; (o) TBSCl, imidazole, 0-23 °C, 3 h, 82%; (p) Fmoc-Ala-Cl, AgCN, PhMe, 23 °C, 18 h, 14%; (q) HF–EtOH (1:4), 0 °C, 1 h, 97%; (r) Jones' reagent, acetone, -15 to 0 °C, 3 h.

chemical yield remained to be improved. Deprotection of the silyl ether in 14 and oxidation of the resulting 15 with Jones' reagent afforded ester segment 16.

As depicted in Scheme 2, the amide segment 24 was prepared from the known α -methylcysteine 17, which was converted to thiazoline 19 in four steps according to the Kelly method¹⁴ for preparation of a thiazoline from cysteine. S-Tritylation of 17 with trityl alcohol and borontrifluoride etherate, esterification of the protected amino acid with thionyl chloride and methanol, condensation of the methyl ester with Fmoc-(S)-Ala-OH and EDCI/DIEA, and deprotective cyclization of the resulting dipeptide 18 with triphenylphosphine oxide and triflic anhydride afforded thiazoline 19 in 53% yield from 17. Deprotection of the Fmoc group in 19 was achieved by use of TBAF in THF-DMF and reprotection with Boc₂O followed by saponification with lithium hydroxide provided thiazoline building block 20. Coupling of 20 with N-MeOHIle 22 derived from 21,⁶ which was previously prepared from the bicyclic α , β -unsaturated lactam,¹⁵ was difficult by even using several coupling reagents due to the severe steric congestion. Fortunately, use of BMTB/DIEA^{11,16} successfully effected the condensation to give amide 23 in 94% yield. However, the thus-obtained 23 was found to be a mixture of the diastereomers. Careful experiment revealed that the saponification of Boc-(S)-(ala)Thz-OMe with lithium hydroxide caused severe epimerization at the α -position of the thiazoline. The mixture was used for further elaboration because the problem could be solved at the later step. Deprotection of the Boc group with TMSOTf and 2,6-lutidine provided amide segment 24.

Activation of ester segment 16 to acid chloride 25 with oxalyl chloride in methylene chloride followed by coupling with 24 in the presence of DIEA smoothly proceeded to afford a mixture of linear precursor 26a and epimer 26b at the α -position of the thiazoline in 89% yield (Scheme 3). Fortunately, protection-free precursor 27a from the respective deprotection of the Fmoc group and the benzyl ester with diethylamine and Pd black was separable with epimer 27b by chromatography and was obtained in 40% yield along with 35% yield of 27b. Macrolactamization of pure 27a using HATU¹¹/DIEA furnished protected halipeptin A (28a) in 61% yield together with 29% yield of epimer 28b, which was generated from racemization at the *N*-MeOHIle residue



Scheme 2. Synthesis of the amide segment 24. Reagents and conditions: (a) TrOH, BF₃·OEt₂, AcOH, 80 °C, 0.5 h; (b) SOCl₂, MeOH, reflux, 11 h; (c) Fmoc-Ala-OH, EDC, DIEA, CH₂Cl₂, 0-23 °C, 10.5 h, 70% (three steps); (d) Ph₃P(O), Tf₂O, CH₂Cl₂, -18 to -10 °C, 3.5 h, 75%; (e) TBAF, THF–DMF (5:1), 0-23 °C, 5 h; (f) Boc₂O, 23 °C, 2 h, 87% (two steps); (g) LiOH, THF–H₂O (4:1), 0-23 °C, 1.5 h; (h) TMSOTf, 2,6-lutidine, CH₂Cl₂, 90%; (i) BMTB, DIEA, CH₂Cl₂, 0-23 °C, 94% (from 22); (j) TMSOTf, 2,6-lutidine, CH₂Cl₂, 23 °C, 2 h, 77%.



Scheme 3. Construction of halipeptin A. Reagents and conditions: (a) $(COCl)_2$, CH_2Cl_2 , 23 °C, 5 h; (b) 24, DIEA, CH_2Cl_2 , 0–23 °C, 6.5 h, 89% from 15; (c) Et_2NH –MeCN (1:1), 0–23 °C, 1.5 h, 97%; (d) H_2 , Pd-black, MeOH–buffer (10:1), 23 °C, 8 h, 40% and 35% of the epimer; (e) HATU, DIEA, CH_2Cl_2 , 23 °C, 24 h, 61% and 29% of the epimer; (f) 4 M HCl–dioxane, ether, 0–23 °C, 4 h, 35% and 50% of the epimer.

during the cyclization. Final deprotection of the TBS group in **28a** was carried out using 4 M HCl–dioxane, which caused again extensive epimerization at the α -position of the thiazoline. Pure halipeptin A (1)¹⁷ was obtained after chromatographic purification in 35% yield. Interestingly, when [(*R*)-(ala)Thz]-epimer **27b** was subjected to macrolactamization under the same conditions, the TBS ether of [(*R*)-(ala)Thz]-halipeptin A was obtained in 73% yield without any epimerization at the *N*-MeOHIle residue.

In conclusion, we have succeeded in total synthesis of halipeptin A through stereoselective construction of the HTMMD fragment. Further studies on the synthesis of halipeptins and the relatives for biological evaluation are underway.

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- Abbreviations: DMP (Dess-Martin periodinane), DEAD (diethyl azodicarboxylate), KHMDS (potassium hexamethylsilazide), EDCI (1-(3-dimethylaminopropyl)-3ethylcarbodiimide hydrochloride), DMAP (4-(dimethylamino)pyridine), DIEA (N,N-diisopropylethylamine), BMTB (2-bromo-3-methyl-4-methylthiazolium bromide), HATU (O-(7-azabenzotriazol-1yl)-N,N,N',N'-tetramethyluronium hexafluorophosphate).
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- 17. Synthetic halipeptin A: $[\alpha]_{D}^{24}$ –11.8 (*c* 0.035, CHCl₃) (lit. –16.6 (*c* 0.029, CHCl₃)); IR v_{max} 3851, 3451, 2925, 2055, 1749, 1636, 1507, 1455, 1260, 1032, 801, 655 cm⁻¹; ¹H NMR (500 MHz, CDCl₃): δ 0.81 (3H, d, J = 7.0 Hz), 0.91 (3H, t, J = 7.0 Hz), 0.98 (3H, d, J = 6.4 Hz), 1.13 (3H, s), 1.20 (3H, s), 1.25–1.58 (10H, m), 1.42 (3H, d,

 $J = 7.3 \text{ Hz}, 1.48 (3H, s), 1.51 (3H, d, J = 7.0 \text{ Hz}), 1.91-1.92 (1H, m), 2.51-2.52 (1H, m), 2.82 (3H, s), 3.08-3.10 (1H, m), 3.298 (3H, s), 3.302 (1H, d, J = 11.9 \text{ Hz}), 3.63-3.66 (1H, m), 3.77-3.79 (1H, m), 4.16 (1H, d, J = 12.2 \text{ Hz}), 4.71 (1H, d, J = 2.8 \text{ Hz}), 4.76-4.87 (2H, m), 5.08 (1H, d, J = 10.4 \text{ Hz}), 7.01 (1H, J = 8.2 \text{ Hz}), 7.22 (1H, d, J = 7.9 \text{ Hz}); ¹³C NMR (125 MHz, CDCl_3): <math>\delta$ 14.3, 14.4, 18.0, 18.4, 18.5, 22.0, 22.3, 23.1, 26.2, 28.2, 30.7, 31.2, 31.9, 34.2, 35.2, 35.7, 44.3, 45.8, 48.6, 49.6, 56.5, 60.9, 64.7, 80.6, 82.6, 83.9, 169.2, 169.6, 172.5, 173.6, 177.3. FAB-HRMS calcd for C₃₁H₅₅N₄O₇S (M+H): 627.3791. Found: 627.3743.