

Total Synthesis and Assignment of Stereochemistry of Raocyclamide Cyclopeptides from Cyanobacterium Oscillatoria raoi

David J Freeman and Gerald Pattenden*

Chemistry Department, Nottingham University, Nottingham NG7 2RD

Received 29 January 1998; revised 18 February 1998; accepted 19 February 1998

Abstract: Total synthesis demonstrates that the structures and stereochemistries of the cyclopeptides raccyclamide A and B isolated from a cyanobacterium should be altered to those shown in formulae 13 and 14 respectively. © 1998 Elsevier Science Ltd. All rights reserved.

A large number of cyclopeptides have now been isolated from nature whose structures are constituted from "unnatural" *D*-amino acids and/or from unusual, highly modified, amino acids, *eg* thiazole, thiazoline, oxazole and oxazoline aminoalkyl carboxylic acids.¹ These interesting structural features raise a number of questions: *eg* Why has nature engineered the synthesis of *D*-amino acids to elaborate certain cyclopeptides? Is this feature associated with some specific folding of a peptide chain prior to macrocycle formation? Are nitrogen, sulphur and oxygen heteroatoms in the heterocyclic-based amino acid residues involved as ionophoric groups to conformationally constrain the cyclopeptide structures and their biological precursors? Is metal encapsulation and transport² a feature of their formation (*ie* acting as a template) and their *modus operandi* in vivo? In order to answer some of these questions we have selected the lissoclinium, *eg* lissoclinamide 4, 1,³ and the raocyclamide, *eg* raocyclamide A, 2⁴, families of cyclopeptides for special study. We selected these two families since their member show all the "unusual" structural features mentioned above, and their synthesis should be fairly straightforward thereby allowing the effects of modifications in the stereochemistries of their constituent amino acids on *eg* peptide folding, case of macrocyclisation, ionophoric properties, to be evaluated. As a prelude to these detailed investigations we have investigated a total synthesis of the structure 2 proposed for raocyclamide A.

Raocyclamide A 2, together with raocyclamide B 7, were isolated from the cyanobacterium *Oscillatoria raoi* ⁴ and their structures and stereochemistries were determined by detailed nmr analysis in combination with amino acid analysis following ozonolysis and application of Marfey's HPLC method. Both molecules show hexapeptide structures formally derived from *D*-isoleucine, *D*-phenylalanine, *D*-serine, and *L*-alanine with their oxazole and thiazole moieties originating from serine and cysteine respectively.

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Our straightforward strategy for the synthesis of raccyclamide A 2, proceeding *via* raccyclamide B 7, and relied on access to the *L*-alanine derived 2,4-disubstituted oxazole 3 and the *D*-isoleucine derived 2,4-disubstituted thiazole 9. We next planned to elaborate 3 to the tetrapeptide 4, then fuse-on the thiazole 9 leading

Reagents: i, AcCl, MeOH, 0°, 1h; ii, Boc-D-Phe-D-Ser-OH, EDC, HOBt, Hunig's base, THF, 0° to RT, 18h; iii, acid of **9**, EDC, HOBt, THF, 0° to RT, 18h; iv, LiOH, H₂O, THF, RT, 1.5h; v, TFA:CH₂Cl₂ (1:1), RT, 1h;vi, DPPA, Hunig's base, THF, high dilution, 4d; vii, Burgess' rgt, THF, 67°, 0.5h.

Scheme 1

to 5, and finally cyclise 6 to raccyclamide B 7, en route to 2. We and others have earlier synthesised the *L*-alanine oxazole 3, in connection with other studies.⁵ After removing the BOC protecting group in 3, a coupling reaction between the corresponding free amine and BOC-*D*-Phe-*D*-Ser-OH then gave rise to the tetrapeptide 4 in 94% yield (Scheme 1).⁶ The substituted thiazole 9 was derived from BOC-protected *D*-isoleucine 8 in six steps (overall yield 34%) as illustrated in Scheme 2.⁷ Saponification of the ester group in 9, followed by a coupling of the resulting carboxylic acid with the amine derived from 4, next led to the hexapeptide 5. After sequential removal of the amine and carboxylic acid protecting groups in 5,

Reagents: i, DCC, HOBt, NMM, CH_2Cl_2 , 0° to RT, 18h; ii, TBSCI, Imidazole, DMF, 0° , 18h; iii, Lawesson's rgt, PhH, 80°, 18h; iv, TBAF, THF, RT, 1.5h; v, Burgess' rgt, THF, 66°, 0.5h; vi, BrCCl₃, DBU, CH_2Cl_2 , 0° to RT, 18h.

macrocyclisation of 6 in the presence of Hünig's base and diphenylphosphoryl azide (DPPA) then produced the cyclic peptide 7. Finally, treatment of 7 with Burgess' reagent⁸ provided the structure 2 proposed for natural raccyclamide A from O. raci. Unfortunately, when we compared the pmr and cmr spectra for our synthetic materials 7 and 2, with those recorded for naturally derived raccyclamides A and B, there were considerable differences. We cogitated the problem and analysed the nmr data in more detail and came to the conclusion that the natural raccyclamides A and B were most likely derived from L-isoleucine rather than its D-enantiomer. Accordingly, we synthesised the disubstituted thiazole 10 from L-isoleucine, then elaborated 10 to 11, and coupled the free carboxylic acid produced from 11 to the free amine derived from 3 leading to 12. Deprotection of the ester and amino groups in 12, followed by macrolactamisation then produced the cyclopeptide 13, which was smoothly cyclodehydrated

Reagents: i, LiOH, H₂O, THF, RT, 1.5h; ii, HCl.NH₂-D-Phe-D-Ser-OMe, EDC, HOBt, Hunig's base, THF, 0° to RT, 18h; iii, Amine of 3, EDC, HOBt, THF, 0° to RT, 18h; iv, TFA:CH₂Cl₂ (1:1), RT, 1h; v, DPPA, Hunig's base, THF, high dilution, 4d; vi, Burgess' rgt, THF, 66°, 0.5h.

Scheme 3

using Burgess' reagent to the oxazoline-based cyclopeptide 14. Gratifyingly, the pmr and cmr spectra for synthetic 13 and 14¹⁰ were identical in every detail with those recorded for natural raccyclamides A and B respectively. Further studies are now in progress to evaluate the effects of stereochemistry of the constituent amino acids and of metal binding, on peptide folding and case of ring closure leading to diastereomers of the raccyclamides 13 and 14, and selected analogues.

Acknowledgements: We thank The Lawes Trust for a scholarship (to DJF) and Jeffrey Hannam for his contribution to this study. We also thank Dr S Carmeli for providing nmr spectra recorded for natural raocyclamide A, and raocyclamide B.

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- P.m.r. data for synthetic 7: δ_H (360 MHz, CDCl₃) 0.88 (3H, d, *J* 6.8, C*H*₃CHCH), 0.99 (3H, t, *J* 7.3, C*H*₃CH₂), 1.25 (1H, m, CH₃CH₂), 1.51 (3H, d, *J* 7.1, C*H*₃CHNH), 1.55 (1H, m, CH₃CH₂), 2.04 (1H, m, (CH₃)CH(C₂H₅)), 3.09 (1H, dd, *J* 7.8 13.7, CH₂Ph), 3.20 (1H, dd, *J* 5.6, 13.7, CH₂Ph), 3.45 (1H, br. s, CH₂OH), 3.67 (1H, dd, *J* 4.2, 10.9, CH₂O), 3.97 (1H, m, CH₂O), 4.43 (1H, m, CHCH₂O), 5.01 (1H, m, CHCH₂Ph), 5.23 (1H, dd, *J* 5.1, 7.7, NHCHCH), 5.33 (1H, m, NHCHCH₃), 7.21 (5H, m, *Ph*), 7.78 (1H, s, SCH), 7.83 (1H, d, *J* 7.2, NHCO), 7.84 (1H, d, *J* 8.0, NHCO), 8.18 (1H, s, OCH), 8.32 (2H, m, 2 x NHCO).
- 10. P.m.r. data for synthetic **14**: δ_{II} (360 MHz, CDCl₃) 0.95 (3H, t, *J* 7.4, C*H*₃CH₂), 1.06 (3H, d, *J* 6.8, C*H*₃CHCH), 1.20 (1H, m, C*H*₂CH₃), 1.40 (1H, m, C*H*₂CH₃), 2.04 (1H, m, (CH₃)C*H*(C₂H₅)), 3.13 (1H, dd, *J* 3.3 12.8, C*H*₂Ph), 3.15 (1H, dd, *J* 2.8, 12.8, C*H*₂Ph), 4.56 (1H, dd, *J* 8.7, 10.7, C*H*₂O), 4.65 (2H, m, C*H*CH₂O, CC*H*CH₃), 4.94 (1H, dd, *J* 6.3, 8.7, C*H*₂O), 5.27 (1H, m, C*H*CH₂Ph), 5.56 (1H, dd, *J* 3.7, 8.8, NHC*H*CH), 6.66 (1H, tt, *J* 1.4, 7.2, CHCHC*H*), 6.81 (4H, m, Ph), 7.34 (1H, d, *J* 4.6, N*H*CHCH₃), 8.04 (1H, d, *J* 8.9, N*H*CHCH₂Ph), 8.20 (1H, d, *J* 8.7, N*H*CHCH), 8.22 (1H, s, SC*H*), 8.22 (1H, s, OC*H*); δ_C (90.5 MHz, CDCl₃) 11.9 (q), 15.7 (q), 20.2 (q), 24.2 (q), 37.7 (t), 42.3 (d), 44.8 (d), 48.0 (d), 55.6 (d), 67.6 (d), 69.9 (d), 124.6 (d), 125.9 (d), 127.6 (d), 129.5 (d), 134.9 (s), 135.6 (s), 141.0 (d), 147.8 (s), 159.3 (s), 159.5 (s), 164.1 (s), 167.0 (s), 169.5 (s), 170.1 (s).