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Asymmetric syntheses of enantiopure C(5)-substituted transpentacins *via* diastereoselective Ireland–Claisen rearrangements[†]

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Asymmetric syntheses of (S,S,S)-2-amino-5-methylcyclopentanecarboxylic acid and (S,S,S)-2-amino-5-phenylcyclopentanecarboxylic acid were achieved in 9 steps from commercially available starting materials *via* the Ireland–Claisen rearrangement of two enantiopure β -amino allyl esters, followed by ring-closing metathesis, reduction and deprotection.

There has been considerable interest in the secondary structural characteristics of oligomers of the cyclic β-amino acids cispentacin and transpentacin (the enantiopure diastereoisomers of 2-aminocyclopentanecarboxylic acid).^{1,2} In order to enhance the structural diversity of enantiopure monomeric cispentacin and transpentacin derivatives available we have developed efficient parallel kinetic resolution (PKR) procedures for the asymmetric syntheses of C(3)- and C(5)-substituted analogues 1-3,³ and subsequently investigated the secondary structural characteristics of some of their oligomers.⁴ Herein we report an alternative procedure for the syntheses of 1,2-anti-1,5-syn-diastereoisomers 6, which are not accessible using the PKR protocol. It was predicted that the Ireland-Claisen rearrangement⁵ of silyl ketene acetals derived from β-amino allyl esters 4 would proceed via a chair-like transition state 7 in which 1,3-allylic strain between the C(3)-substituents and the C(1)-O bond is minimised, and rearrangement occurs on the face opposite the bulky N-benzyl-N-(amethylbenzyl) group. The resultant α -substituted β -amino acid products 5 could then be elaborated to the corresponding C(5)-substituted transpentacins 6 via ring-closing metathesis and hydrogenolytic reduction/deprotection (Fig. 1).

Rearrangement precursors **11** and **12** were prepared from commercially available sorbic acid **8** in four steps. Esterification of **8** upon treatment with isobutylene/H⁺ gave α , β -unsaturated ester **9** in 80% isolated yield. Conjugate addition of lithium (*S*)-*N*-benzyl-*N*-(α methylbenzyl)amide⁶ to **9** produced the known β -amino ester **10**^{7,8} as a single diastereoisomer (>99:1 dr), which was isolated in 89%



Fig. 1 Substituted cispentacin and transpentacin analogues 1–3 and the Ireland– Claisen rearrangement strategy to access transpentacins 6.

yield and >99:1 dr after chromatographic purification. Hydrolysis of the tert-butyl ester moiety within 10 upon treatment with TFA, conversion of the resultant carboxylic acid to the corresponding acid chloride, and treatment with either alcohol 23 or alcohol 24 (which were both prepared in >99:1 dr upon reduction of the corresponding alkynes 21 and 22) gave β -amino allyl esters 11 and 12 in 57 and 63% yield, respectively, over the two step procedure. Ireland-Claisen rearrangement of 12 was achieved via deprotonation with LiHMDS and treatment of the resultant lithium (E)- β -amino enolate⁹ with TMSCl, followed by heating the corresponding silvl ketene acetal at reflux in PhMe, which gave a 76:24 mixture of β -amino acids 15 and 16, respectively. Conversion of this mixture to the corresponding methyl esters 19 and 20 (for ease of handling and isolation) was then achieved upon treatment with MeI and DBU,¹⁰ and 19 and 20 were then isolated as single diastereoisomers (>99:1 dr) in 69 and 16% yield, respectively (Scheme 1). The relative configurations

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within both **19**·HBF₄ and **20** were unambiguously assigned by single crystal X-ray diffraction analyses (Fig. 2),¹¹ with the absolute configurations within (*S*,*S*,*S*,*S*,*E*)**·19** and (2*R*,3*S*,1′*R*, α *S*,*E*)**·20** being assigned from the known configuration of the (*S*)- α -methylbenzyl fragment. Analogous Ireland–Claisen rearrangement of **11** produced a (separable) 80:20 mixture of β -amino acids **13** and **14**, which after esterification gave **17** and **18** in >99:1 dr and 45 and 14% isolated yield, respectively (Scheme 1). The configurations within both **17** and **18** were initially assigned by analogy to those within **19** and **20**, and these assignments were subsequently confirmed by chemical correlation and single crystal X-ray diffraction analysis of a derivative (*vide infra*). In both cases, the configurations within the major diastereoisomers **17** and **19** were consistent with the predicted stereochemical outcome of transition state model **7** (Fig. 1).

The configuration within **18** (the minor diastereoisomer derived from rearrangement of **11**) was unambiguously established by chemical correlation: ring-closing metathesis of **18** (>99:1 dr) upon treatment with Grubbs I catalyst gave **25** in 77% yield and >99:1 dr. Subsequent hydrogenolytic reduction of **25** in the presence of Pd/C gave a complex mixture of products including the known β -amino ester **26**, which was isolated in 9% yield after exhaustive purification



Fig. 2 X-ray crystal structures of (S,S,S,S,E)-**19**·HBF₄ [left] and $(2R,3S,1'R,\alpha S,E)$ -**20** [right] (selected H atoms are omitted for clarity).

of the crude reaction mixture by flash column chromatography (Scheme 2). The ¹H and ¹³C NMR data for this sample of **26** were in excellent agreement with those reported previously;^{3d} this analysis therefore enabled the configurations within **18**, **25** and **26** to be assigned.

For both 17 and 19 (the major diastereoisomers arising from Ireland-Claisen rearrangement of 11 and 12, respectively), ringclosing metathesis in the presence of Grubbs I catalyst gave 27 and 28 as single diastereoisomers (>99:1 dr) in 85 and 80% yield,¹² respectively, and subsequent tandem hydrogenation/hydrogenolysis gave primary amines 29 and 30 as single diastereoisomers (>99:1 dr) in 83 and 87% yield (Scheme 3). The relative configurations within 27-30 were initially assigned by ¹H NMR nOe analyses and these assignments were subsequently confirmed unambiguously by single crystal X-ray diffraction analyses (Fig. 3):¹¹ in the case of 29, single crystal X-ray diffraction analysis of the corresponding 2,6-difluorobenzoic acid salt and the determination of a Flack xparameter¹³ of 0.005(11) for this crystal structure allowed the assigned absolute (S,S,S)-configuration within 29 to be confirmed, whilst for 30, single crystal X-ray diffraction analysis of the corresponding HBF₄ salt and the determination of a Flack *x* parameter¹³ of -0.03(16) for the crystal structure of 30 HBF₄ allowed the assigned absolute (S,S,S)-configuration within 30 to also be confirmed.









Fig. 3 X-ray crystal structures of (*S*,*S*,*S*)-**29**·2,6-difluorobenzoic acid·CHCl₃ [left] and (*S*,*S*,*S*)-**30**·HBF₄ [right] (selected H atoms and CHCl₃ are omitted for clarity).

These crystallographic studies therefore also secured the assigned configurations within **17**, **27** and **28**. Finally hydrolysis of the methyl ester functionalities within both **29** and **30**, upon treatment with 6.0 M aq HCl at reflux for 16 h, gave β -amino acids **31** and **32** which were isolated as single diastereoisomers (>99:1 dr) in 84 and 89% yield, respectively, after purification on Dowex 50WX8 ion exchange resin (Scheme 3).

In conclusion, the Ireland–Claisen rearrangement of two enantiopure β -amino allyl esters, followed by ring-closing metathesis, reduction and deprotection, enabled the asymmetric syntheses of (*S*,*S*,*S*)-2-amino-5-methylcyclopentanecarboxylic acid and (*S*,*S*,*S*)-2amino-5-phenylcyclopentanecarboxylic acid in 9 steps and >99:1 dr in both cases, and 18 and 19% overall yield, respectively, from commercially available starting materials.

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- 11 X-ray crystal structure data were collected using an Oxford Diffraction SuperNova diffractometer with graphite monochromated Cu-Ka radiation, using standard procedures at 150 K. The structures were solved by direct methods (SIR92); all non-hydrogen atoms were refined with anisotropic thermal parameters. Hydrogen atoms were added at idealised positions. X-ray crystal structure data for 19 HBF₄ [C₃₁H₃₆BF₄NO₂]: M = 541.43, orthorhombic, $P2_12_12_1$, a = 9.3325(1) Å, b = 15.3112(2) Å, c = 20.0967(2) Å, V = 2871.65(5) Å³, Z = 4, $\mu = 0.784$ mm⁻¹, colourless block, crystal dimensions = $0.23 \times 0.26 \times 0.30 \text{ mm}^3$. A total of 6010 unique reflections were measured for 4 $< \theta <$ 77 and 5068 reflections were used in the refinement. The final parameters were $wR_2 = 0.102$ and $R_1 = 0.038 [I > -3.0\sigma(I)]$. CCDC 936128.[†] X-ray crystal structure data for 20 $[C_{31}H_{35}NO_2]$: M = 453.62, monoclinic, P_{21} , a = 7.9925(2) Å, b =16.7779(4) Å, c = 10.6442(3) Å, $\beta = 111.582(3)^\circ$, V = 1327.29(7) Å³, Z = 2, $\mu =$ 0.541 mm^{-1} , colourless plate, crystal dimensions = 0.05 \times 0.20 \times 0.24 mm^3 . A total of 4653 unique reflections were measured for 4 < θ < 77 and 4635 reflections were used in the refinement. The final parameters were $wR_2 = 0.083$ and $R_1 = 0.034 [I > -3.0\sigma(I)]$. CCDC 936129.⁺ X-ray crystal structure data for 29 $C_7H_4F_2O_2$ CHCl₃ [$C_{16}H_{20}Cl_3F_2NO_4$]: M =434.69, monoclinic, C2, a = 25.0016(5) Å, b = 6.6915(1) Å, c = 12.9605(2) Å, $\beta = 110.830(2)^\circ$, V = 2026.55(7) Å³, Z = 4, $\mu = 4.452$ mm⁻¹, colourless prism, crystal dimensions = $0.11 \times 0.12 \times 0.29 \text{ mm}^3$. A total of 4226 unique reflections were measured for 4 $< \theta <$ 76 and 4207 reflections were used in the refinement. The final parameters were $wR_2 = 0.082$ and $R_1 = 0.033 [I > -3.0\sigma(I)]$. CCDC 936130.[†] X-ray crystal structure data for **30** HBF₄ [C₁₃H₁₈BF₄NO₂]: M = 307.09, monoclinic, $P2_1$, a = 5.5150(2) Å, b =7.6423(3) Å, c = 17.2563(6) Å, $\beta = 90.531(3)^{\circ}$, V = 727.27(5) Å³, Z = 2, $\mu = 1.099 \text{ mm}^{-1}$, colourless block, crystal dimensions = $0.22 \times 0.23 \times 0.23$ 0.27 mm³. A total of 2045 unique reflections were measured for 5 < θ < 76 and 2038 reflections were used in the refinement. The final parameters were $wR_2 = 0.092$ and $R_1 = 0.036 [I > -3.0\sigma(I)]$. CCDC 936131⁺.
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