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Selective Epimerisation of a Fungal Cyclopeptolide via an 2-Amino-oxazole Intermediate - Conformational Consequences

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Abstract

The selective epimerisation of the cyclic depsipeptide (cyclopeptolide HUN-7293) **1** at a single amino acid (Leu⁵) was achieved in four steps. A *regio*-selective thionation at one of its six amide-bonds provided the corresponding thioamide **2**, which upon *S*-benzylation followed by an unusual silver-promoted *intra*-molecular cyclisation, was converted to the 2-amino-oxazole derivative **4**. Re-opening of the oxazole ring by acid hydrolysis regenerated the cyclopeptolide, but preferentially with inverted stereochemistry at Leu⁵ **5**. The influence of this inversion on the backbone conformation is discussed.

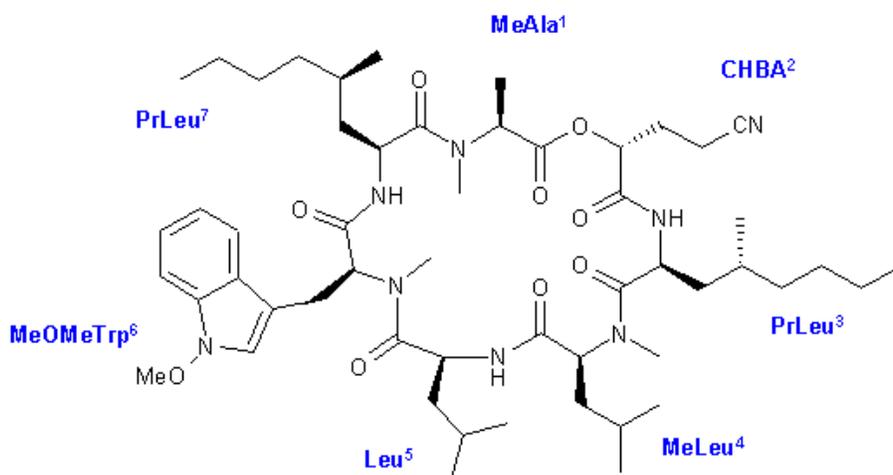


Fig. 1: Structure of the fungal metabolite HUN-7293 (**1**)

Introduction

The cyclic depsipeptide HUN-7293 **1**, a fungal fermentation product, potently inhibits VCAM-1 expression on activated endothelial cells^[1]. In addition to *N*-methylated amino acids MeAla and MeLeu, it contains the uncommon constituents D-4-cyano-2-hydroxybutyric acid (CHAB), *N*1'-methoxy-*N*-methyltryptophan (MeOMeTrp) and d-propylleucine (PrLeu). Its constitution (Fig. 1) and 3-dimensional structure has been determined by 2D-NMR methods and X-ray diffraction analysis (Fig. 2)^[2]. The 3-dimensional structure is characterised by two methylated *cis*-amide bonds and two *trans*-annular hydrogen bonds resulting in a tightly packed backbone conformation. This allows a close hydrophobic contact of the sidechains of hydroxy acid² (CHBA²), amino acid⁶ (MeOMeTrp⁶) and amino acid⁵ (Leu⁵) as indicated by a strong magnetic shielding effect of the aromatic ring on the Leu⁵ b-protons (Leu⁵ b-H: $\delta = -0.36$ ppm).

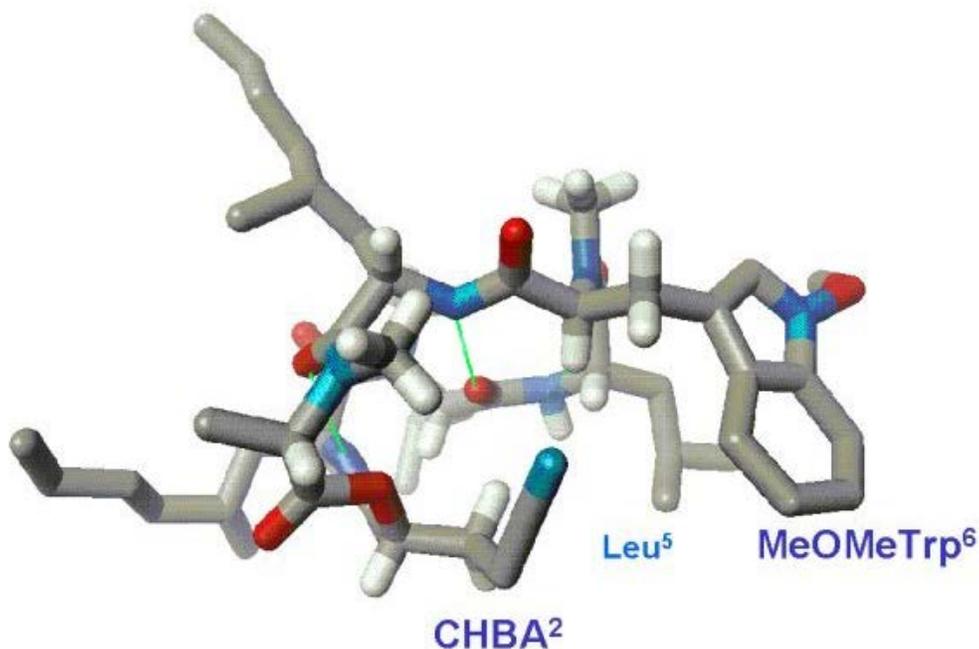
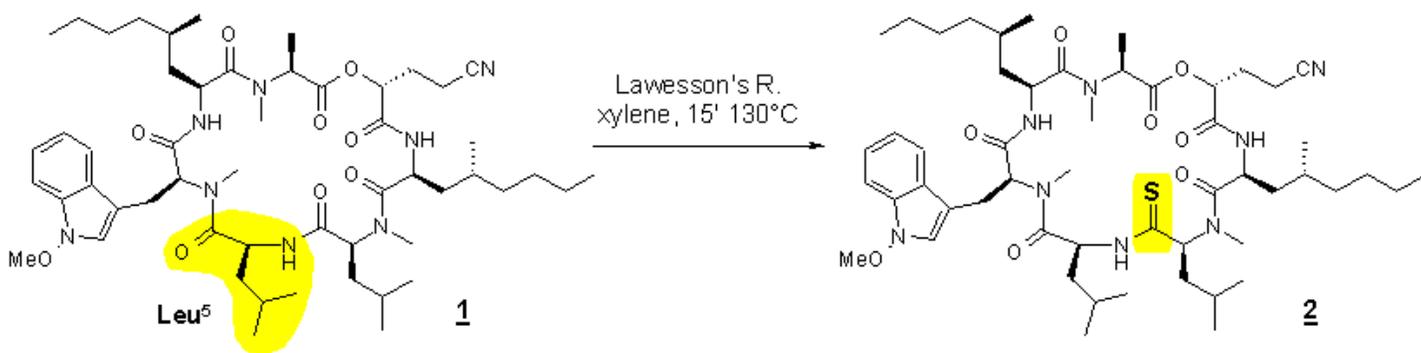


Fig. 2: X-ray crystal structure of HUN-7293, 3D: [msv-file](#) or [mol-file](#)

Results and Discussion

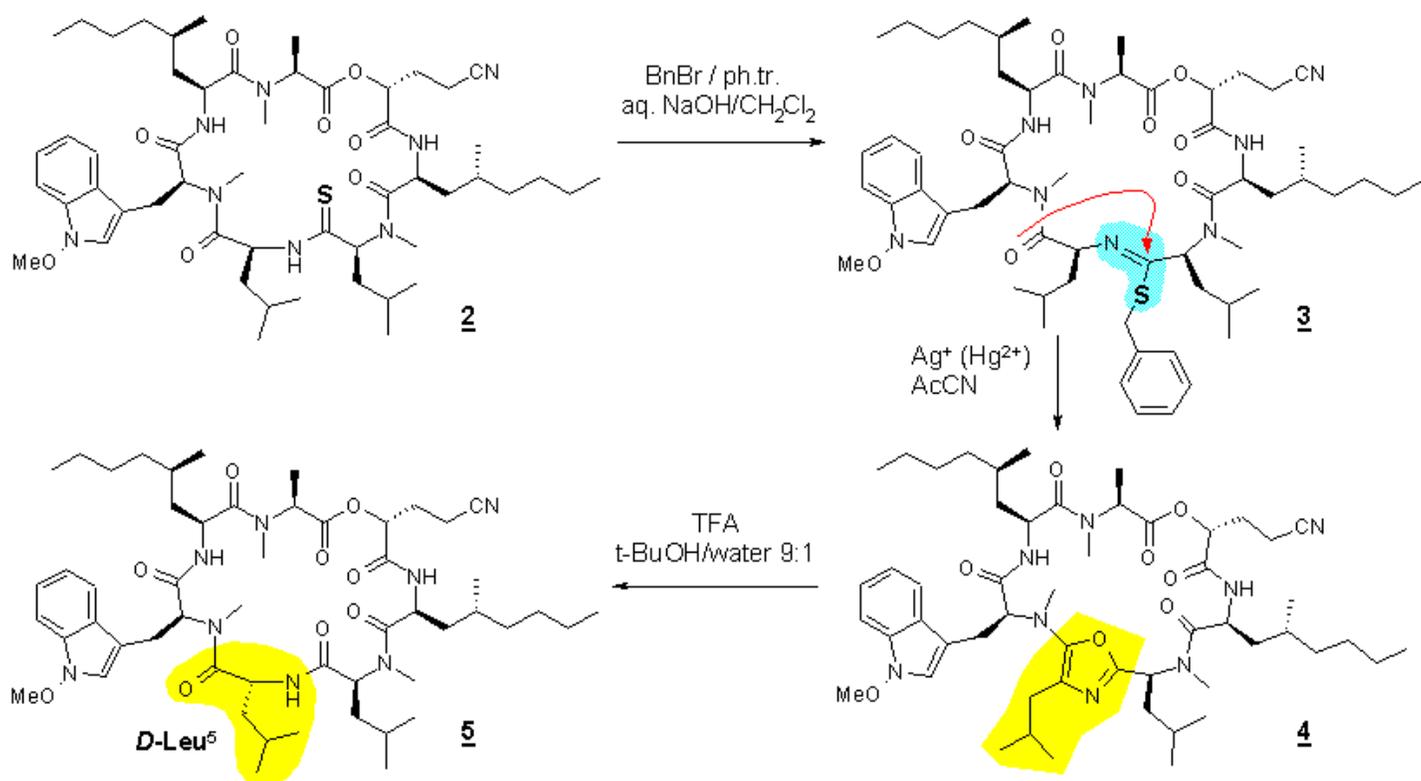
Highly *regio*-selective thionation of the most reactive amide carbonyl between MeLeu⁴ and Leu⁵ (the only primary amide NH-group, which is not involved in *intra*-molecular hydrogen bonding) was easily achieved with 0.5 equivalents Lawesson's reagent in xylene at 130 °C (Scheme 1). As by-products, the double thioamide (additional thionation at MeOMeTrp⁶-PrLeu⁷) and a triple thioamide (thionation at all primary amides) could be isolated.



Scheme 1: Selective transformation of a single amide bond with Lawesson's reagent

Alkylation of the mono-thioamide **2** with benzyl bromide under phase-transfer conditions (aq. NaOH, dichloromethane, ultrasound) resulted in a quantitative transformation into the S-benzyl thioamidate **3**. Treatment of this thioamidate with silver- or mercury salts in acetonitrile destroyed the chirality at Leu⁵ by the unusual formation of a peptide-bridged 2-aminoxazole **4**. In contrast to other oxazole-containing peptidic natural products, no sidechain atoms are involved in the ring-closure of this rather exotic type of a rigid dipeptide mimetic^[5-9]. The formation of the heterocycle can be explained via the activation of the thioamidate by co-ordination of the heavy metal at the sulphur (i.e. formation of an excellent leaving group). Due to the lack of an external nucleophile, this forces the molecule to undergo an *intra*-molecular cyclisation

involving the adjacent carbonyl group of Leu ([Scheme 2](#)).



Scheme 2: 2-Aminooxazole as achiral dipeptide-mimetic

Utilising the sensitivity of 2-aminooxazoles towards acid hydrolysis, the two masked amid bonds could easily be regenerated by treatment with TFA in aqueous *t*-butanol. This led to the formation of the $D\text{-Leu}^5$ epimer **5** together with HUN-7293 itself in a 2:1 ratio. The predominance of one diastereomer probably reflects the rigid structure of the peptide-bridged oxazole, as also indicated by a single NMR-conformation. The stereochemistry of the predominant diastereomer **5** was established by 2D-NMR methods and independently proven via partial synthesis.

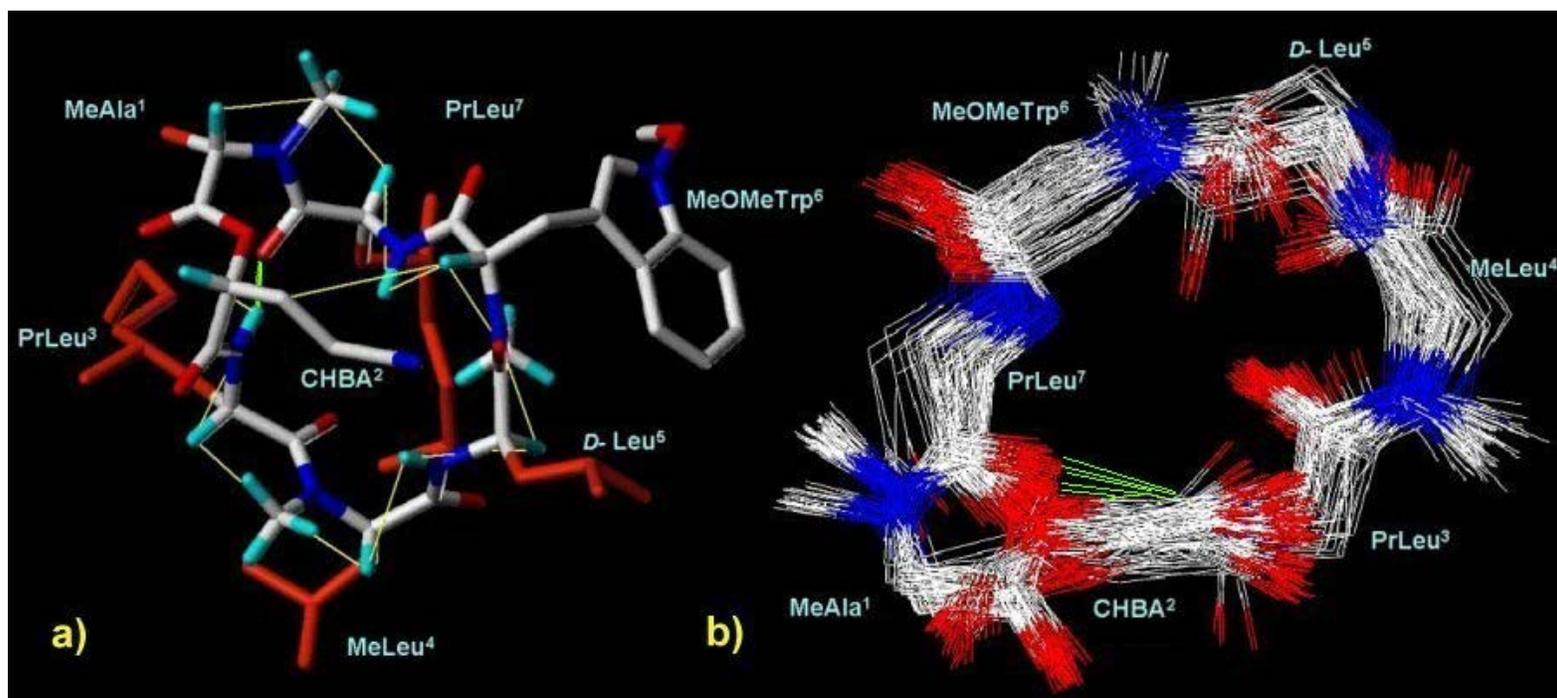


Figure 3:

a) Observed ROESY distance constraints (yellow) in an low energy conformation with a possible hydrogen bridge (green) PrLeu³ NH - CO PrLeu⁷ (3D: [msv-file](#) or [mol-file](#))

b) Ensemble of 100 energy-minimised backbone structures compatible with the distance constraints (simulated annealing^[10]); Due to the lack of distance constraints at the lactone moiety, two equally populated conformations at MeAla¹ are visible.

In contrast to HUN-7293 **1**, the 5-*epi* derivative **5** adopts a single conformation in CDCl₃ solution. Although dramatic changes in the conformation could be expected due to the epimerisation, 5-*epi* HUN-7293 shares some characteristic NMR-features with the parent compound:

- high field shift of the Leu⁵ NH
- ROESY cross-peaks of CHBA²-MeOMeTrp⁶ indicating a "back-folding" of the CN-sidechain onto the ring ([Fig. 3a](#))

The main difference was the absence of the extreme high-field shift of one Leu⁵ b-proton. A simulated annealing analysis was performed utilising the available set of backbone ROESY distance-constraints ([Fig. 3b](#)). The resulting structure was compatible with the observed ³J_{NH-Ha} coupling constants, whereas coupling constants of sidechain protons ³J_{Ha-Hb} could only be explained by a mixed population of sidechain conformers.

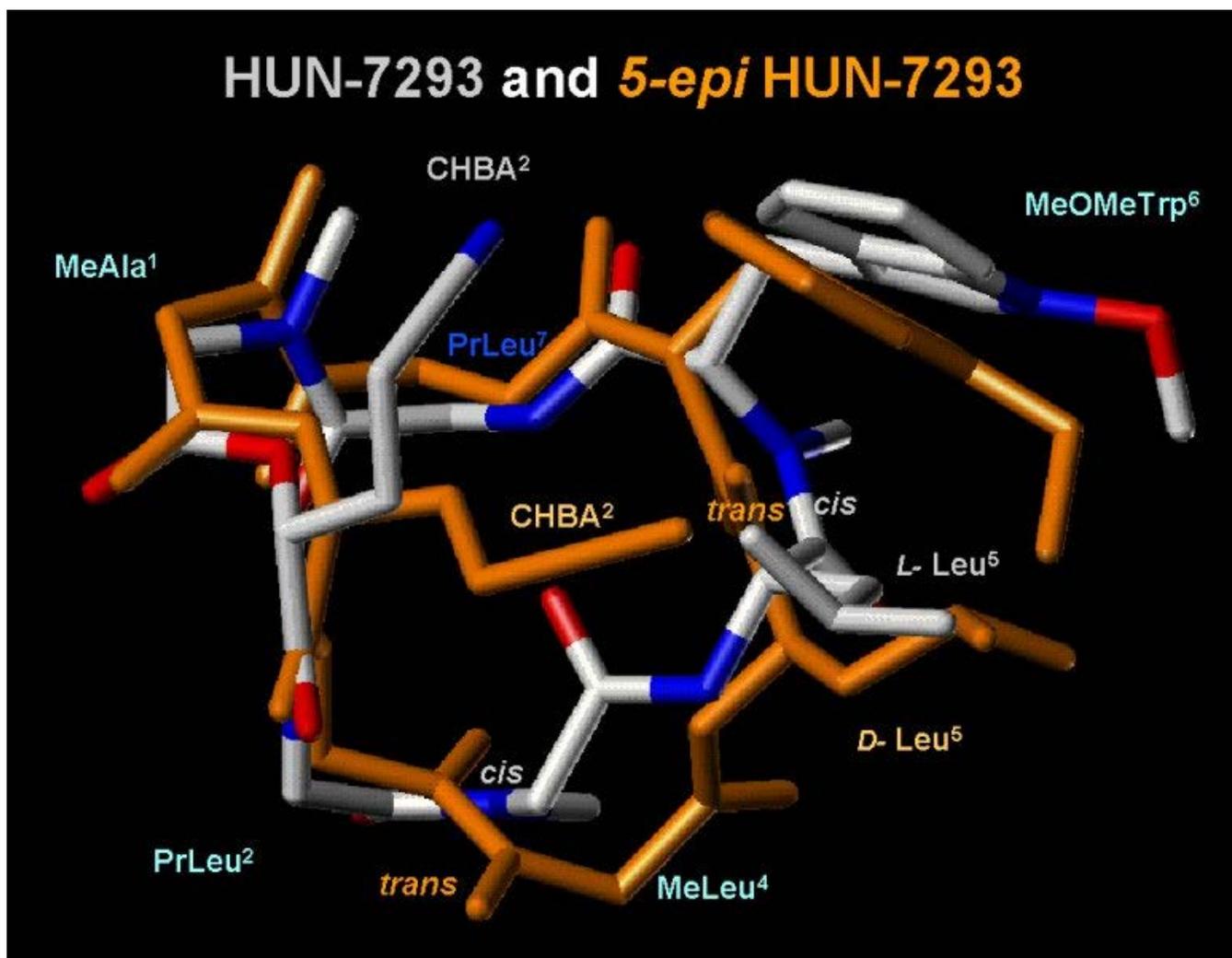


Figure 4: Overlay of the natural product HUN-7293 (X-ray structure, grey) and its Leu⁵-epimer (NMR structure, orange; 3D: [msv-file](#) or [mol-file](#))

Conclusions

Because of the known sensitivity of the cyclopeptolides 3D conformation towards even minor structural changes, we were surprised to see that the *5-epi* HUN-7293 adopts an overall similar conformation. The different chirality of Leu⁵ is obviously compensated by the flipping of both *cis*-amide bonds to the *trans*-conformation still allowing at least one *trans*-annular hydrogen bond and the hydrophobic interaction of the CHBA², MeOMeTrp⁶ and Leu⁵ (Fig. 4).

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 10. simulated annealing: Sybyl, 100 cycles 700K 1000fs, 200K 1000fs, energy minimisation: Tripos forcefield, Gasteiger charges, no violations of distance constraints greater than 20 pm (ROESY)
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Comments

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