

Total Synthesis and Structural Confirmation of the Marine Natural Product Dysinosin A: A Novel Inhibitor of Thrombin and Factor VIIa

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The marine natural product dysinosin A 1¹ is a new member of serine protease inhibitors generally known as the aeruginosins (Figure 1).² It exhibits activity against thrombin, an essential enzyme in the blood coagulation cascade,³ and Factor VIIa which is involved in blood vessel damage in complex with tissue factor (TF).⁴ The structure of dysinosin A was determined by detailed NMR studies and its absolute stereochemistry deduced from an X-ray structure of a complex with thrombin.¹ Dysinosin A possesses unique structural and functional features that distinguish it among the aeruginosins. Noteworthy is the presence of a 5*S*,6*R*-dihydroxy octahydroindole carboxylic acid, an unusual guanidine as part of a pyrrolidine ring,^{2d,5} and a sulfate group. Bonjoch,⁶ Wipf,⁷ and their respective groups have independently reported the total synthesis and stereochemical revision of aeruginosin 298-A utilizing L-tyrosine as a starting material.

We report herein the total synthesis of dysinosin A utilizing a carbon construct strategy that generates subunits originating from L-glutamic acid, butyrolactone, D-leucine, and D-mannitol as shown in Figure 1. The synthesis of the enantiopure octahydroindole carboxylic acid⁸ capitalized on the prospects of a ring-closure metathesis reaction⁹ from a chiron derived from L-pyroglutamic acid, and subsequent stereoselective epoxidation and epoxide opening. To this end we had to secure methodology that introduced two C-allylic appendages with a *syn*-disposition at C-4 and C-5 of L-proline as shown in Scheme 1. The (4*S*)-allyl analogues **2** and **3** have been previously synthesized by stereoselective enolate alkylation of the corresponding L-glutamate esters.¹⁰ Conversion of **2** and **3** into the corresponding methyl L-pyroglutamates,¹¹ selective reduction, and acetylation afforded the expected hemiaminal derivatives **4** and **5**. The introduction of a *syn*-allyl group at C-5 via *N*-acyl iminium ion chemistry¹² proved to be challenging. After extensive variations of solvents, the nature of Lewis acids, and *N*-substituents,¹³ allylation of **5** could be realized with a 5.5:1 all-*syn/anti* selectivity with allyl tributylstannane in the presence of BF₃·Et₂O in toluene to afford **7**, easily separable from the minor diastereomer by chromatography. Allylation of **4** under the same conditions led to a 1:2 ratio of *syn/anti* isomers of **6**.

Olefin metathesis of **6** and **7** using the original and elegant Grubbs method⁹ led to the carbocyclization products **8** and **9**, respectively, in excellent yields. Epoxidation with *m*-CPBA proved to be highly selective, affording the epoxides **10** and **11** in each case, presumably as a result of an attack from the more accessible convex face of the bicyclic system. When treated with aqueous TFA, each epoxide led to the enantiopure intermediates **12** and **13**, respectively, whose structures were unequivocally proven by single-crystal X-ray analysis. For reasons of functional group compatibility, the synthesis was continued with **13**, which was transformed to

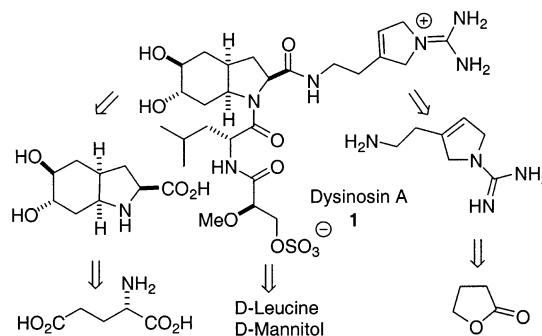
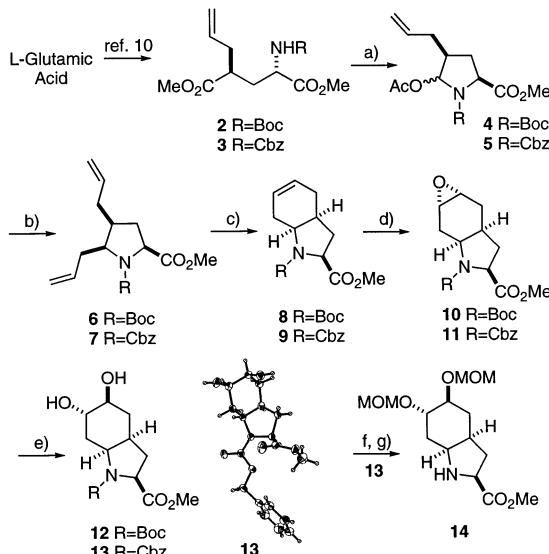


Figure 1. Disconnection of dysinosin A to subunits and chirons.

Scheme 1^a

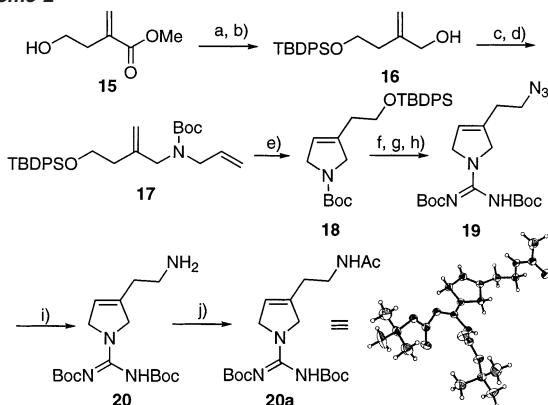


^a Reagents and conditions: (a) 1. TFA, CH₂Cl₂; 2. NaHCO₃; 3. Δ, toluene; 4. LiHMDS, CbzCl, THF -78 °C; 5. LiHBET₃, THF -78 °C; 6. Ac₂O, DMAP, CH₂Cl₂; overall 85%. (b) BF₃·OEt₂, allyl tributylstannane, toluene -78 °C (*syn/anti* 5.5:1); overall 83%. (c) Ru benzylidene(Cy₃P)₂Cl₂ 1 mol %, CH₂Cl₂; 99%. (d) *m*-CPBA, CH₂Cl₂. (e) TFA (0.2 equiv), THF/H₂O (1/1); 75–79% (2 steps). (f) MOMCl, (iPr)₂NEt, CH₂Cl₂; 98%. (g) Pd/C 20%, H₂, MeOH; 95%.

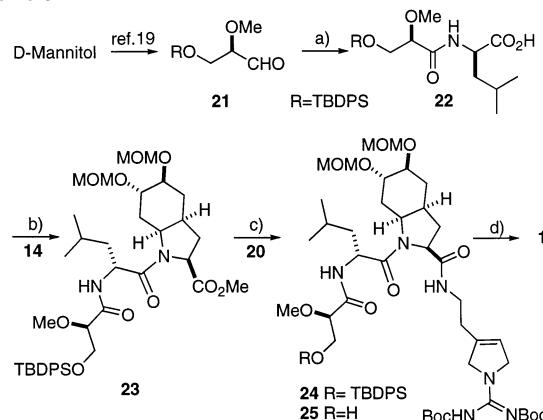
the bis-MOM ether **14**. The highly site-selective *trans*-dialixal acid-catalyzed opening could be due in part to the shielding effect of the pseudodioxial¹⁴ carbomethoxy group on the concave face of the bicyclic motif, as evidenced by X-ray analysis (Scheme 2).

The synthesis of the Δ-3 pyrrolidine unit¹⁵ shown in Scheme 2, started with the hydroxy ester **15** readily available from butyrolactone.¹⁶ Reduction of the ester function gave the allylic alcohol **16**,¹⁷ which was further transformed to the diolefin **17** in high overall yield. The versatility of the Grubbs metathesis reaction⁹ and its tolerance of functional groups was evidenced by the

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Scheme 2^a

^a Reagents and conditions: (a) TBDPSCl, imidazole, DMF; 90%. (b) DIBAL-H, CH₂Cl₂; 90%. (c) MsCl, Et₃N, CH₂Cl₂; then allylamine; 84%. (d) Boc₂O, Et₃N, CH₂Cl₂; quant. (e) Ru benzylidene(Cy₃P)₂Cl₂ 10 mol %, CH₂Cl₂; 90%. (f) TBAF, THF; 92%. (g) PPh₃, DEAD, (PhO)₂P(ONa), THF; 82%. (h) TFA, CH₂Cl₂; then Et₃N, Goodman's reagent; 86%. (i) PPh₃, H₂O, THF then AcOH; 72%. (j) Ac₂O, Et₃N, MeOH; 90%.

Scheme 3^a

^a Reagents and conditions: (a) 1. NaClO₂, 2-methylbut-2-ene, NaH₂PO₄, t-BuOH; 2. TFA-d-Leu-Bn, EDC, HOEt, CH₂Cl₂; 3. Pd/C 10%, H₂, MeOH; overall 76%. (b) 14, BopCl, (iPr)₂NEt, MeCN; 63%. (c) 1. LiOH, THF/MeOH; 2. EDC, HOEt, CH₂Cl₂; overall 92%. (d) 1. TBAF, THF; 2. Py-SO₃⁻, Bu₂SnO, CH₂Cl₂; 6 h; 3. TFA, CH₂Cl₂; 6 h; prep. HPLC; 34% overall.

successful cyclization of **17** to the pyrroline **18** in 90% yield.¹⁸ A series of well-precedented transformations gave **20** which was definitively characterized by single-crystal X-ray analysis of the corresponding *N*-acetyl derivative **20a**.

The acyclic peptide chain **22** was prepared as shown in Scheme 3 from d-leucine and 2-*O*-methyl-d-glyceraldehyde easily available from d-mannitol.¹⁹ Peptide coupling between **14** and **22** afforded **23** which was hydrolyzed to the free acid. A second peptide coupling with **20** proceeded in good overall yield to give **24**, which was desilylated to the alcohol **25**. Treatment of **25** with pyridine-SO₃ complex in dichloromethane in the presence of a catalytic quantity of dibutyltin oxide²⁰ afforded the corresponding sulfate ester. Hydrolysis of the *N*-Boc group with TFA in dichloromethane, followed by isolation of the crude product and purification by reverse phase HPLC afforded dysinosin A as a white solid, identical in all respects to the natural product (HPLC, ¹H, ¹³C NMR, FAB and electrospray mass spectrometry).

The total synthesis of dysinosin A by an enantioselective route provides definitive evidence for its structural and configurational identity. It also represents the first total synthesis of a hitherto unknown member of the aeruginosin family of marine antithrombin natural products, with inhibitory activity against Factor VIIa.

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Supporting Information Available: Experimental procedures of key reactions, NMR, and X-ray and other data (PDF). This material is available free of charge via the Internet at <http://pubs.acs.org>.

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