

The Synthesis of Functionalized Nanoscale Molecular Rods of Defined Length

Christopher G. Levins and Christian E. Schafmeister*

Department of Chemistry, University of Pittsburgh, Pittsburgh, Pennsylvania 15260

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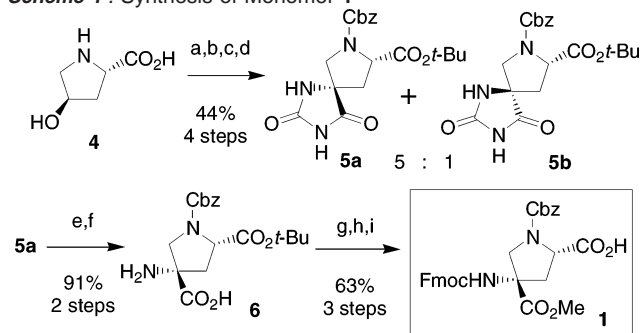
The development of systematic approaches to the synthesis of nanometer scale molecules of discrete size, shape, and mechanical properties is a long-term goal of macromolecular chemistry.^{1–10} Molecular rods are macromolecules that have many exciting potential applications in nanoscale science as spacers, wires, and construction elements; yet as Michl's authoritative review¹¹ points out, the poor solubility of many current examples of molecular rods is the primary obstacle to their study and application. We have developed a synthetic approach to water-soluble molecular rods that are of defined length and contain multiple chiral centers. These molecules can be easily synthesized and functionalized on both ends. We present here the synthesis of a prototype molecular building block **1** and demonstrate the solid-phase synthesis of two

molecular rods **2** and **3** containing three and five monomers, respectively. Molecular mechanics was used to predict the conformation of the scaffold **2**, and the prediction was confirmed by 2D NMR analysis.

Our prototype building block **1** was synthesized on a 1.8 g scale from inexpensive, commercially available *trans*-4-hydroxy-L-proline **4** in nine steps with an overall yield of 25% (Scheme 1). The synthesis uses a Bucherer–Bergs reaction¹² to install a quaternary stereocenter with a diastereoselectivity of 5:1.¹³ Using a single chromatographic column, we isolated 16.5 g of pure diastereomer **5a** from a 20 g mixture of **5a** and **5b**. The intermediate **5a** leads to the differentially protected bis amino acid **1**.

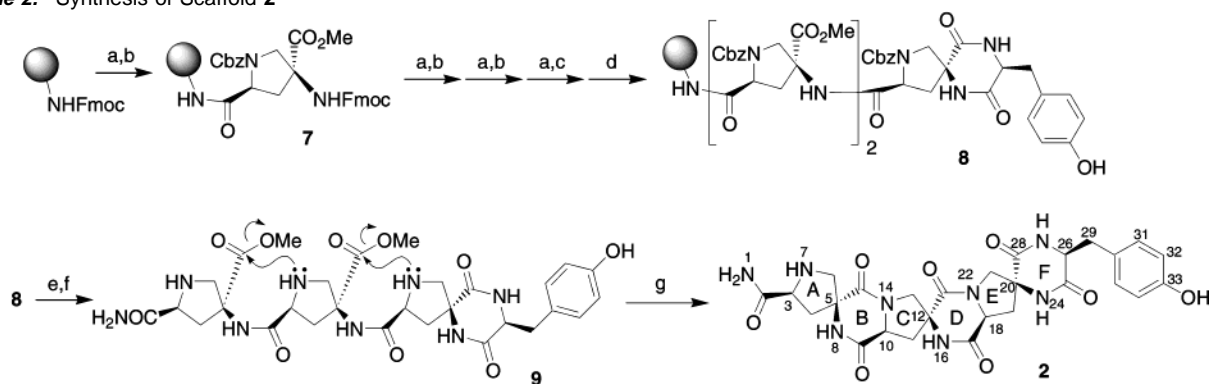
We assembled three building blocks to form the molecular rod **2** using sequential solid-phase synthesis on a 46 μ mol scale (Scheme 2). Each building block was activated as the 1-hydroxy-7-azabenzotriazole (HOAt) ester,¹⁴ and quantitative coupling to the previous building block was achieved in less than 10 min, a surprising result given the apparent hindered nature of the nucleophile. After three monomers were coupled, the oligomer was functionalized with a tyrosine residue to facilitate characterization. The flexible oligomer **8** was cleaved from the resin, and high-performance liquid chromatography with mass spectrometry (HPLC-MS) analysis revealed a single peak with the expected *m/z* ratio of 1061.2 (*M* + *H*⁺). The carboxybenzyl (Cbz) groups were then removed by hydrogenolysis to form **9**. The flexible oligomer **9** was converted into the rigidified scaffold **2** through the parallel formation of two diketopiperazine¹⁵ rings by exposure to 20% piperidine/DMF over 48 h at 4 °C.¹⁶ The product **2** precipitated from the 20% piperidine solution, and filtration was used to recover 5 mg (~15% yield based on resin loading). HPLC-MS analysis of this material reveals a single peak with the expected *m/z* ratio of 595.2 (*M* + *H*⁺) for **2**. The scaffold **2** is soluble in water to more than 5 mg/mL and stable in neutral and acidic aqueous solution at room temperature for weeks.

Scheme 1. Synthesis of Monomer 1^a



^a Reagents and conditions: (a) (i) TMS–Cl, NEt(*i*Pr)₂, CH₂Cl₂, reflux; (ii) Cbz–Cl, 0 °C to room temperature. (b) Jones reagent, acetone, 20 °C. (c) Isobutylene, H₂SO₄ (cat.), CH₂Cl₂, room temperature. (d) (i) (NH₄)₂CO₃, KCN, 1:1 DMF/H₂O, 60 °C, sealed tube; (ii) silica gel chromatography, 99:1 CH₂Cl₂/MeOH. (e) (Boc)₂O, DMAP, THF, room temperature. (f) KOH, 1:1 H₂O/THF, room temperature. (g) (i) TMS–Cl, NEt(*i*Pr)₂, CH₂Cl₂, reflux; (ii) Fmoc–Cl, 0 °C to room temperature. (h) (i) MeOH, DCC, DMAP, CH₂Cl₂, 0 °C to room temperature; (ii) silica gel chromatography 2:1 EtOAc/hexanes. (i) CF₃CO₂H/CH₂Cl₂.

Scheme 2. Synthesis of Scaffold 2^a



^a Reagents and conditions: (a) 20% piperidine/DMF, 40 min, room temperature. (b) Two times 2 equiv of **1**–OAt ester, 4 equiv of NEt(*i*Pr)₂, 20% CH₂Cl₂/DMF, 60 min, room temperature. (c) 3 equiv of *N*- α -Fmoc-*O*-*tert*-butyl-L-Tyr-OAt ester, 6 equiv of NEt(*i*Pr)₂, 20% CH₂Cl₂/DMF, 60 min, room temperature. (d) 20% piperidine/DMF, 120 min, room temperature. (e) 95% CF₃CO₂H/2.5% (*i*Pr)₃SiH/2.5% H₂O, 120 min, room temperature. (f) H₂ (~1 atm), 10% Pd/carbon, 7:2:1 MeOH/H₂O/AcOH, 60 min, room temperature. (g) 20% piperidine/DMF, 48 h, 4 °C.

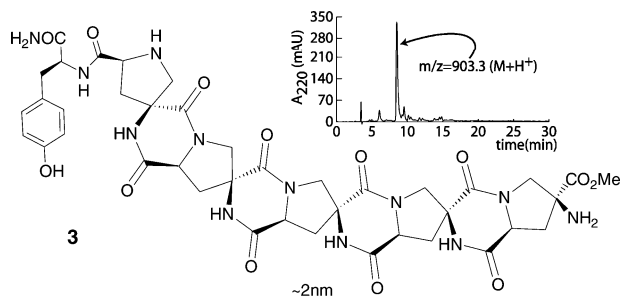


Figure 1. The chemical structure of scaffold **3**. Modeling indicates that the spiro-fused ring structure forms a narrow left-handed helical rod with approximately four residues per turn and a pitch of ~ 20 Å. Inset: the C_{18} reverse phase HPLC chromatogram of the unpurified ether precipitated product containing **3** (0–20% MeCN/H₂O, 0.1% TFA).

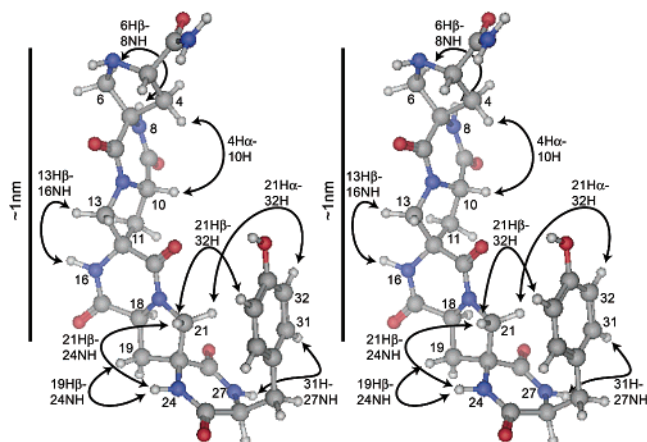


Figure 2. A stereoimage of the predicted third lowest energy conformer of scaffold **2** most consistent with NMR data. The arrows indicate that strong and medium ROESY cross-peaks are seen between the indicated pairs of protons. The molecule forms a molecular rod about 1.5 nm in length.

To demonstrate the generality of this synthetic approach, the five-mer scaffold **3** (Figure 1) was synthesized in a fashion similar to that of scaffold **2**. The resin (49 mg, 31.4 μmol loading) was first charged with an Fmoc-protected tyrosine residue, and then five cycles of coupling using monomer **1** were performed. Roughly 13 mg of resin was removed and subjected to cleavage conditions. The Cbz groups were removed and the scaffold was rigidified by exposure to 20% piperidine/DMF over 24 h. In this case, about 3 mg of the scaffold **3** ($\sim 33\%$ from initial resin loading) was isolated by precipitation with ether and centrifugation. After all of these manipulations, this unpurified material was remarkably homogeneous (Figure 1), and HPLC-MS analysis confirmed that the major peak has the expected m/z ratio of 903.3 ($M + H^+$). This material was soluble in 10% D₂O/H₂O at 5 mg/mL.

To construct a model of the scaffold **2**, we carried out an *in vacuo* conformational search using the AMBER95 force field¹⁷ within the molecular mechanics package MOE.¹⁸ The conformational search revealed a cluster of five lowest energy conformations all within 0.4 kcal/mol of each other separated from the next highest energy cluster by a gap of 2.2 kcal/mol. This energy gap predicts that the molecule will spend roughly 98% of its time collectively in the five lowest energy minima (Figure 2). A superposition of these predicted lowest energy conformations reveals that rings B,

C, D, E, F and the folded tyrosine conformation are identical. The differences between the five conformations involve combinations of rotamers around the C2–C3 bond, rotamers around the tyrosine OH bond, and two envelope conformations of ring A. The 2D ROESY spectra display cross-peaks consistent with those from the predicted rigid B–F ring system. The NMR data are more consistent with ring A existing predominately in the single envelope conformation depicted in Figure 2 based upon a strong cross-peak between 4H α and 10H and a weak or nonexistent cross-peak between 4H β and 10H.

We have demonstrated the synthesis of two nanoscale molecular rods using a novel building block approach. The synthesis takes place in two phases: a flexible chain is first assembled through amide coupling on solid support, and then the scaffold is rigidified by the selective formation of a second set of amide bonds. We have also demonstrated that the three-mer scaffold has a rodlike three-dimensional structure. The scaffolds were rapidly assembled using solid-phase synthesis and were easily functionalized on either of their two ends. Among molecular rods, these scaffolds exhibit the rare property of water solubility,¹¹ making them compatible with biological buffers. Because of their fused ring structure, we expect them to be more inflexible than spacers such as poly-proline helices.^{19,20} This synthetic approach may also be extended to incorporate other building blocks to construct more complex shapes and ultimately functional macromolecules.

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Supporting Information Available: Synthesis and characterization of **1**, **2**, and **3**. HPLC-MS and 2D NMR data for **2** and HPLC-MS data for **3** (PDF). This material is available free of charge via the Internet at <http://pubs.acs.org>.

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