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Synthesis of a Series of Polar, Orthogonally Protected, α,α -Disubstituted Amino Acids

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Abstract: The synthesis of four novel, "cationic", $\alpha_i \alpha_i$ -disubstituted amino acids is described. The new amino acids use an orthogonal protection scheme making them suitable for incorporation via solid-phase peptide synthesis. © 1997 Elsevier Science Ltd.

 α,α -Disubstituted amino acids ($\alpha\alpha$ AAs), **1** (where R, R' \neq H), are widely used in peptide design because of their structure promoting effects.¹ α -Aminoisobutyric acid (Aib) and Aib-like residues (where R, R' are not bulky) have been incorporated into peptides to form 3₁₀ or α -helices depending upon the design,² percentage of $\alpha\alpha$ AAs,³ and location of the $\alpha\alpha$ AAs.⁴ Unfortunately, most $\alpha\alpha$ AAs that have been incorporated are hydrophobic which precludes spectroscopic or crystallization experiments of peptides containing high percentages of $\alpha\alpha$ AAs in aqueous media.^{1,2b} Very few examples of polar $\alpha\alpha$ AAs suitable for incorporation into peptides have been reported in the literature.^{5,6} Polar $\alpha\alpha$ AAs are a must for the synthesis of short, highly helical, water soluble peptides containing high percentages of $\alpha\alpha$ AAs. These peptides are integral to the study of the 3₁₀/ α -helix equilibrium, the investigation of the stability of the 3₁₀-helix in aqueous media, and in the design of short antimicrobial peptides.^{2,7} Herein we report the synthesis of a series of polar $\alpha\alpha$ AAs suitable for incorporation into peptides are integral to the study of the 3₁₀/ α -helix equilibrium, the peptides.^{2,7} Herein



The general structure of the amino acids is shown above, **2**. The amino acids are constructed from a 6membered ring backbone due to the high helix promoting effects previously exhibited by cyclic $\alpha\alpha$ AAs in short peptides.^{2a,7} The polar functionality is introduced via reductive amination which provides flexibility for the incorporation of a wide variety of R" groups, such as hydrophobic chains, additional polar groups, or fluorescent probes. This side chain nitrogen will be protonated under physiological conditions and under conditions normally used for aqueous spectroscopic characterization.

The synthesis begins with a reductive amination on the commercially available 1,4cyclohexanedione monoethylene ketal **3** with the amine of choice (1.1 equiv.), acetic acid (1.0 equiv.), and sodium triacetoxyborohydride (1.6 equiv.) in 1,2-dichloroethane to give **4**.⁸ Sodium cyanoborohydride was also tried as the reducing agent but gave unsatisfactory yields as compared to the acetoxy compound. The ketone functionality is unmasked by treating **4** with a 20% trifluoroacetic acid/water solution and heating to reflux in THF for 24 hours to yield **5**. To obtain an amino acid suitable for Fmoc solid-phase peptide synthesis, the side chain must be orthogonally protected with respect to the α -nitrogen. Thus, the nitrogen on ketone **5** is protected with *t*-butyloxycarbonyl (Boc) using Et₃N (0.95 equiv.), (Boc)₂-O (1.14 equiv.) and a catalytic amount of DMAP in THF to give **6**.



Ketone **6** is then converted to hydantoin **7** using a Bucherer-Bergs procedure.^{5,9} At this point, conventionally the hydantoin would be hydrolyzed using harsh conditions (i.e. barium hydroxide in a ParrTM Bomb) which could effect the Boc protecting group. Therefore, we use a mild hydrolysis method developed by Rebek in which the hydantoin nitrogens are activated by reaction with (Boc)₂-O to give **8**.^{5,10} The selective hydrolysis is accomplished with 1N lithium hydroxide (8 equiv..) and THF as a co-solvent at room temperature to give amino acid **9**. The α -nitrogen is protected with fluorenylmethyloxycarbonyl (Fmoc) using TMS-Cl (2.5 equiv.), DIEA (3.0 equiv.), and Fmoc-Cl (1.1 equiv.) in methylene chloride. Protecting the α -nitrogen of a very hydrophobic amino acid of this type is difficult using traditional methods with aqueous/organic mixtures because of the extreme non-polar nature of the side chain (protected). The method we use, developed by Bolin and co-workers,¹¹ solubilizes the amino acid by forming the silyl ester in neat organic media thereby allowing protection of hydrophobic amino acids. The crude product obtained from the Fmoc reaction is purified over silica gel using CHCl₃/MeOH mixtures to give pure **2**.¹²



The method provides a general route to polar $\alpha\alpha AAs$ with R groups of varying hydrophobicity. This method could also be easily adapted to the synthesis of fluorescent or other tagged $\alpha\alpha AAs$. We plan to study the effects of the hydrophobic chains on antimicrobial activity and helicity.

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- 12. **2a**; 1-amino(N α -Fmoc)-4-(N δ -Boc, N δ -ethyl)cyclohexane-1-carboxylic acid. ¹H NMR (250 MHz, DMSO d₆) δ 7.94 (d, 2H), 7.79 (d, 2H), 7.62 (s, 1H), 7.45 (t, 2H), 7.38 (t, 2H), 4.32 (m, 3H), 3.82 (bm, 1H), 3.16 (bm, 2H), 2.25 (bm, 2H), 1.74 (bm, 4H), 1.50 (bm, 2H), 1.46 (s, 9H), 1.12 (t, 3H). ¹³C NMR (100 MHz, DMSO *d*₆) δ 175.88, 155.24, 154.24, 143.82, 140.73, 127.65, 127.00, 125.33, 120.01, 78.26, 65.20, 57.28, 54.20, 46.75, 43.80, 31.02, 28.15, 25.05, 16.02. FAB-MS (glycerol) m/z 509.3 (M+H)⁺. 15.1% overall yield. 2b; 1-amino(N^{α}-Fmoc)-4-(N^{δ}-Boc, N^{δ} -butyl)cyclohexane-1-carboxylic acid. ¹H NMR (300 MHz, DMSO d₆) δ 7.88 (d, 2H), 7.73 (d, 2H), 7.56 (s, 1H), 7.42 (t, 2H), 7.31 (t, 2H), 4.26 (m, 3H), 3.70 (bm, 1H), 3.02 (bm, 2H), 2.16 (bm, 2H), 1.69 (bm, 4H), 1.42 (bm, 4H), 1.40 (s, 9H), 1.24 (m, 2H), 0.89 (t, 3H). ¹³C NMR (50MHz, CDCl₃) & 178.54, 155.68, 155.38, 143.73, 141.28, 127.64, 127.01, 124.96, 119.89, 79.47, 66.63, 58.16, 54.00, 47.19, 43.56, 32.76, 31.64, 28.47, 25.21, 20.25, 13.88. FAB-MS (glycerol) m/z 537.5 (M+H)⁺. 28.4% overall yield. 2c; 1-amino(N^{α}-Fmoc)-4-(N^{δ}-Boc, N^{δ}benzyl)cyclohexane-1-carboxylic acid. ¹H NMR (250 MHz, CDCl₃) δ 7.73 (d, 2H), 7.57 (d, 2H), 7.36 (t, 2H), 7.29 (m, 7H), 4.89 (s, 1H) 4.41 (d, 2H), 4.33 (s, 2H), 4.16 (t, 1H), 4.00 (bm, 1H), 2.11 (bm, 2H), 1.79 (bm, 2H), 1.54 (bm, 4H), 1.40 (s, 9H). 13 C NMR (100 MHz, DMSO d₆) δ 175.87, 155.40, 155.25, 143.81, 140.72, 140.02, 128.15, 127.63, 127.02, 126.70, 126.49, 125.35, 120.08, 79.19, 65.21, 57.25, 54.16, 46.76, 46.10, 31.02, 28.01, 25.07. FAB-MS (glycerol) m/z 571.2 (M+H)⁺. 13.7% overall yield. 2d; 1-amino(N^{α}-Fmoc)-4-(N^{δ}-Boc, N^{δ}-2naphthylmethyl)cyclohexane-1-carboxylic acid. ¹H NMR (300 MHz, DMSO d_6) δ 8.48 (s, 1H), 7.96 (m, 2H), 7.80 (m, 2H), 7.69 (m, 3H), 7.30 (bm, 8H), 4.92 (s, 2H), 4.85 (d, 1H), 4.75 (m, 2H), 3.60 (bm, 1H), 2.13 (bm, 2H), 1.74 (bm, 4H), 1.51 (bm, 2H)1.46 (s, 9H). ¹³C NMR (100 MHz, CDCl₃) δ 183.93, 155.99, 155.85, 143.86, 141.41, 134.68, 133.79, 130.92, 129.01, 127.75, 127.14, 126.20, 125.99, 125.78, 125.65, 125.40, 125.10, 123.44, 119.99, 80.29, 66.61, 58.20, 54.50, 47.40, 45.00, 32.72, 28.52, 25.55. FAB-MS (glycerol) m/z 621.0 (M+H)⁺. 18.7% overall yield.

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