## Stereospecific synthesis of chiral alkinylogous amino acids

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*tert*-Butoxycarbonyl (Boc)-protected  $\alpha$ -amino aldehydes, conventionally prepared from the corresponding (S)- $\alpha$ -amino acids, are converted *via* the Corey–Fuchs reaction into the *N*-protected alkinylogous amino acids, which on deprotection yield the corresponding enantiomerically pure acids.

The synthesis of unusual amino acids continues to be an interesting endeavour for several reasons,<sup>1</sup> *e.g.* the need to prepare peptidomimetic drugs.<sup>2</sup> In this connection the incorporation of alkinylogous amino acids 1 has not been considered to date. It can be anticipated that the presence of one or more of such units in a peptide chain may influence, *inter alia*, the secondary and tertiary structure. Indeed, preliminary molecular modelling studies show that polypeptidic forms of 1 are expected to be in an  $\alpha$ -helical form, as in the case of normal  $\alpha$ -amino acids, but with a considerably larger inner helix-channel.<sup>3</sup> Here we describe the synthesis and properties of enantiomerically pure alkinylogous amino acids 1.<sup>4</sup>

Our strategy is based on the Corey-Fuchs transformation<sup>5</sup> of Boc-protected  $\alpha$ -amino aldehydes 3<sup>6</sup> derived from the corresponding S-configurated amino acids 2 (Scheme 1). The enantiomeric purity of the crude dibromides 4 was checked by removing the Boc-group and forming the corresponding (R)-Mosher amides.<sup>7</sup> HPLC analysis indicated a diastereoisomeric excess (de) for 4a, b, c and d of 90, 70, >99 and >99%, respectively.8 In the case of the alanine derivative, simple recrystallization of 4a led to a product having 99% ee (40%) yield). The synthesis of the more extensively racemized phenylalanine derivative 4b had to be optimized by carrying out the reaction at -30 °C and recrystallizing the crude product from diethyl ether-hexane (ee = 98%; 82% yield). The use of enantiomerically pure dibromides 4 afforded enantiomerically pure derivatives of alkinylogous amino acids 1. Thus, the Nprotected acids 5 were converted into the corresponding methyl esters by treatment with MeI/K<sub>2</sub>CO<sub>3</sub>/DMF (R = Me, 78%; R =Bn, 66%;  $R = Pr^{i}$ , 82%;  $R = Bu^{i}$ , 80%), which were then deprotected at the nitrogen atom and converted into the corresponding (R)-Mosher amides. HPLC-analyses indicated diastereoisomeric purities of >99%, corresponding to essentially complete enantiomeric purity of the acids 5 (ee  $>99\%).^{8}$ 

The alkinylogous amino acids 1 are white-greyish solids which decompose upon heating to 180–200 °C. In the case of 1a suitable crystals for an X-ray structural analysis were obtained.<sup>†</sup> The results show that a three-dimensional array pertains in which the carboxylate moieties form hydrogen bonds to the NH<sub>3</sub><sup>+</sup>-entities of the neighbours (Fig. 1), the same as in the crystal structure of alanine itself.<sup>9</sup> Thus, the acetylenic spacer not only has the effect of separating the carboxyl moiety from the NH<sub>3</sub><sup>+</sup>-function, it also places the repeating hydrogen bonding units further away from one another in the crystal lattice.



Using the classical titration method, the isoelectric points  $[pK_i = 1/2 (pK_S + pK_B)]$  of the alkinylogous amino acids 1 were determined.<sup>8</sup> The values turned out to be consistently lower than those of the corresponding  $\alpha$ -amino acids (1a, 4.86; 1b, 4.67;



Scheme 1 Reagents and conditions: i, lit.,<sup>6</sup>; ii, CBr<sub>4</sub>–Zn–PPh<sub>3</sub>, 24 h, 0 °C; 3, 2 h, -30 °C; iii, BuLi, 1 h,  $-78 \rightarrow -55$  °C, CO<sub>2</sub>, 2 h,  $\rightarrow 22$  °C; iv, CF<sub>3</sub>CO<sub>2</sub>H–CH<sub>2</sub>Cl<sub>2</sub>, 1 h, 22 °C



Fig. 1 Structure of the alkinylogous alanine 1a in the crystal, showing the hydrogen bonds (symmetry related atoms denoted by \*). Selected interatomic distances (Å) and angles (°): N–C(1) 1.490(2), C(1)–C(2) 1.469(2), C(2)–C(3) 1.189(2), C(3)–C(4) 1.469(2), C(4)–O(1) 1.224(2), C(4)–O(2) 1.244(1), N···O(1)\* 2.718(2), N···O(2)\*\* 2.792(1), N···O(2)\*\*\* 2.754(1), C(1)–C(2)–C(3) 178.1(1), C(2)–C(3)–C(4) 177.6(1), O(1)–C(4)–O(2) 124.2(1), N–C(1)···C(4)–O(2) (torsion) -33.4(4) [alanine N–C–C–O -18.3].

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Scheme 2 Reagents and conditions: i,  $Et_3N$  in THF-H<sub>2</sub>O (1:1), 1 h, 22 °C; ii,  $K_2CO_3$ , DMF, MeI, 22 °C

1c, 5.19; 1d, 4.63 vs. alanine, 6.00; phenylalanine, 5.48; valine, 5.96; leucine, 6.02). This is due to the fact that compounds 1 have lower  $pK_S$  and  $pK_B$  values relative to those of the parent  $\alpha$ -amino acids.

Preliminary attempts to incorporate 1 in dipeptides were successful. Thus, reacting the Boc- or benzyloxycarbonyl (Z)-protected active esters 6 derived from (S)-alanine or (S)-phenylalanine with the protonated form of compounds 1 followed by esterification afforded the N-protected dipeptide esters 7. In a final reaction the diastereoisomeric (R,S) form of 7c was prepared by using (R) phenylalanine and (S) 1d (yield 57%).<sup>8</sup>

Initial biological screening was carried out with some of the alkinylogous amino acids 1. For example, compound 1d was subjected to the Ames test, which showed that it has essentially no mutagenic properties.<sup>‡</sup> However, the biological properties of peptidic forms and of other derivatives of 1 still need to be studied.

## Footnotes

† Crystal data for 1a: C<sub>5</sub>H<sub>7</sub>NO<sub>2</sub>, M = 113.1 g mol<sup>-1</sup>, colourless, 0.25 × 0.25 × 0.42 mm, a = 6.7355(3), b = 7.9237(3), c = 11.6912(6) Å, V = 623.96(5) Å<sup>3</sup>, T = 293 K,  $D_c = 1.20$  g cm<sup>-3</sup>, μ(Cu-Kα) = 0.79 mm<sup>-1</sup>, F(000) = 240 e, Z = 4, orthorhombic, space group  $P2_12_12_1$  [No. 19], Enraf-

Nonius CAD4 diffractometer,  $\lambda = 1.54178$  Å, measuring method  $\omega$ -20, 2676 measured reflections  $(\pm h, \pm k, +l)$ ,  $[(\sin\theta)/\lambda]_{max}$  0.63 Å<sup>-1</sup>, 1278 independent reflections, 1247 observed reflections  $[I \ge 2\sigma(I)]$  for 76 refined parameters, structure solved by direct methods, non H atoms refined anisotropically, H atom positions riding,  $\Sigma w(F_o^2 - F_c^2)^2$  minimized, R = 0.027,  $R_w = 0.081$  [ $w = 1/[\sigma^2(F_o^2) + 0.002p^2 + 0.036p$ ] with  $p = (F_o^2 + 2F_c^2)/3$ , absolute structure determined [Flack 0.0(3)], final difference Fourier  $\Delta = 0.13$  e Å<sup>-3</sup>. Atomic coordinates, bond lengths and angles, and thermal parameters have been deposited at the Cambridge Crystallographic Data Centre. See Information for Authors, Issue No. 1.

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