Synthesis of an L-proline modified mimetic of the A83586C antitumour cyclodepsipeptide[†]

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A mimetic of the A83586C cyclodepsipeptide has been synthesised *via* a three-segment coupling protocol involving dipeptides 9, 8 and 7; in contrast to our previous synthesis of A83586C, where the HATU-mediated macrolactamisation proceeded in 25% yield, the corresponding macro-lactamisation of *seco*-amino acid 18 occurred in *ca*. 78% yield.

A83586C¹ and GE3² are exciting new anticancer agents with a mechanism of action thought to involve the selective inhibition of deregulated E2F transcription factors within cancer cells.²

Some time ago, we reported the first asymmetric total synthesis of A83586C³ and its 4-*epi*-analogue.⁴ 4-*Epi*-A83586C has a (3*R*)-piperazic acid component replacing the (3*S*)-Piz unit. While this modification does actually serve to improve the yield of macrolactamisation from 25 to 70%,^{3,4} it has a seriously detrimental effect on antitumour potency; the 4-*epi*-analogue being much less active as an antitumour drug.⁵ High-field NMR studies on 4-*epi*-A83586C⁴ suggest that its C(8)-carbonyl assumes a *cis*-orientation relative to the C(7)-piperazine-NH, which is opposite to A83586C, where a *trans*-relationship is believed to exist between these two groupings.¹ The combined experimental data suggests that significant conformational perturbations to this region are not well tolerated.



We anticipated obtaining **4** from the chemoselective union of 5^3 with **6**. A [2 + 2 + 2]-fragment condensation was envisaged for constructing the linear hexadepsipeptide precursor of **6**; the key ring-closure would be effected through the L-pro and D-Thr termini (Scheme 1).

Dipeptide 14 (Scheme 2) was prepared from N(Z)-L-proline (10) by amidation with *tert*-butyl carbazate,⁷ hydrogenolytic removal of the Z-group from 11 with a Pd/C catalyst, and silver cyanide mediated coupling⁸ of 12 with 13. Fmoc⁹ cleavage of 14 with Et₂NH in acetonitrile subsequently procured amine 9. The coupling of dipeptides 9 and 8³ proceeded smoothly when mediated by BOP-Cl¹⁰ and Et₃N at low temperature for 1.5 h; tetrapeptide 15 was isolated in 89% yield after chromatography. Fmoc-deprotection and subsequent reaction of 16 with acid chloride 7³ under silver cyanide-assisted conditions⁸ furnished the desired depsipeptide 17 in 73% yield. The Boc-protected



As part of an ongoing programme aimed at identifying a more readily synthesised analogue of A83586C that has similar conformational properties and improved potency, we selected the L-proline-modified congener **4** as a candidate for study. Our choice of an L-proline replacement for the (3S)-piperazic acid

† Electronic supplementary information (ESI) available: characterisation data. See http://www.rsc.org/suppdata/cc/b2/b204018b/



Scheme 1 Retrosynthetic strategy for the L-proline modified A83586C congener (4).

cis to C(7)-NH

10.1039/b20401

ЫÖ



Scheme 2 Synthetic route to the L-proline modified mimetic 6.

acyl-hydrazide of the L-proline residue in **17** was next converted to the corresponding acid by successive treatment with TFA (to remove the Boc groups) and *N*-bromosuccinimide in aqueous THF. The latter reaction is thought to create a hydrolyticallylabile acyl diazene.¹¹ The macrolactamisation of *seco*-aminoacid **18** proceeded smoothly when HATU¹² was used to activate the proline residue under high-dilution conditions. The desired macrolactam **19** was isolated in 48% overall yield for the three steps from **17**. The Troc group of **19** was now detached with Zn in aqueous AcOH¹³ and compound **20** acylated with ZCl to obtain **21**. This protocol enabled **21** to be obtained highly pure by flash chromatography prior to it being hydrogenated over Pd on C in methanolic HCl. The latter process worked efficiently, compound **6** being recovered in essentially quantitative yield.

Future work will attempt to chemoselectively couple **6** to the pyran activated ester 5^3 to obtain **4**. It is envisaged that the solution conformation and antitumour activity of **4** will then be determined to see how they compare with A83586C. Attempts will also be made to exploit cyclodepsipeptide **6** for high-throughput parallel synthesis work aimed to identifying simplified new anticancer drugs that act by an E2F inhibitory mechanism.

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