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Synthesis of N^α-(Purinyl/Pyrimidinyl acetyl)-4-Aminoproline Diastereomers with Potential Use in PNA Synthesis[#]

Bargur P. Gangamani, Vaijayanti A. Kumar* and Krishna N. Ganesh*

Division of Organic Chemistry (Synthesis), National Chemical Laboratory, Pune 411008, INDIA

Abstract: This paper describes an approach to introduce conformational constraint and chirality into the PNA backbone by bridging the ethylenediamine and glycine components of the same unit by a methylene group which leads to PNA based on 4-aminoprolyl backbone with chirality at C-4 and C-2. The synthesis and characterisation of all four diasteroisomers with thymine (T) as the sidechain nucleobase (3a-d) and the synthesis of one of the stereoisomer (2S, 4R) linked to each of the four nucleobases (10-13) are described. Using these monomeric units, two model dimers (17, 18) containing four chiral centres but differing in stereochemistry at only one site were prepared and CD data on these indicate considerable structural differences in base stacking induced by chiral backbone among these.

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INTRODUCTION

Peptide Nucleic Acids (PNAs) are designed structural composites in which the nucleobases of natural DNA (I) are linked to a pseudopeptide backbone (II) of repeating units of N-(2-aminoethyl)glycine.¹ These compounds have recently emerged as novel nucleic acid mimics that efficiently bind DNA/RNA in a sequence specific manner² and hold potential for development as antisense therapeutic agents.³ The extraordinary tight binding of PNA to complementary DNA and RNA has been attributed to a lack of negative charge on PNA, rigidity of the peptide backbone due to the presence of sp^2 hybridization and tertiary amide linkages. Attempts aimed at structural modifications to improve PNA properties such as binding affinity, specificity and cell permeability have centered around chemical modifications of PNA backbone,⁴ and the introduction of asymmetry into achiral PNA⁵⁻⁹ may influence its direction of binding with complementary DNA and RNA. The latter attribute was imparted by (i) covalent linking of a chiral amino acid,⁵ an oligonucleotide⁶ or a polypeptide⁷ at the PNA terminus, (ii) constructing a PNA backbone with an amino acid carrying a nucleobase in the side chain⁸ and (iii) incorporation of various amino acids in PNA backbone itself.⁹

An effective way to explore contribution of backbone rigidity to PNA binding with DNA and RNA, is to employ PNA monomers containing stereospecific chemical modifications. We envisaged the introduction of



chemical modifications into PNA backbone in a way that not only makes them conformationally rigid but also chiral by a strategy as shown in structure III where the α -C of the glycine and β -C of ethylenediamine of thesame monomeric unit are bridged with a methylene group to generate five membered heterocyclic rings along the backbone. The monomer precursors to build such oligomers are 4-(R/S)-amino-N^{α}-(A/G/C/Tylacetyl)-2-(R/S)-prolines IV containing two asymmetric centers at C-4 and C-2. A systematic oligomerisation of all the four possible diastereomers would lead to polychiral PNA having wide conformational diversity in the backbone.⁹ In this paper, we report the preparation of appropriately protected T-monomers of all four diastereomers (**3a-d**) from a single starting material 4R-hydroxy-2S proline (1) and the synthesis of one stereoisomer (2S, 4R) linked to each of the four nucleobases (A, G, C and T) via a common precursor. Further, in order to demonstrate the utility of such monomers, synthesis of two of the derived prolyl-PNA dimers **17** and **18**, each containing four chiral centers but differing in stereochemistry at only one site, is described. Preliminary spectroscopic studies indicate that considerable differences in base stacking are induced by configurational change even at a single site in the backbone.

RESULTS AND DISCUSSION

The naturally occurring 4R-hydroxy-2S-proline (1) was transformed to the designed targets 3a-d by four independent pathways (A-D), each involving a similar set of reactions, but differing in the order of inversion steps at C-4 and C-2 (Scheme-1). Path A consisted of esterification of carboxyl group in 1 and N^{α}protection with BOC, followed by conversion into 4R-tosyl derivative.^{10a} Treatment of this with NaN₃ in DMF resulted in S_N 2 displacement (inversion) of 4R-tosyl group by N₃ to 4S-N₃ derivative^{10b} (2a) that was reduced to 4S-NH₂ compound which was subsequently protected with either CBz or Fmoc group.¹¹ Deprotection of N1 followed by DCC-HOBT coupling with thymin-1-ylacetic acid¹² gave the fully protected 4S,2S T-monomer (3/3a). One of its epimer 2b (4R,2S derivative) was obtained by path B which involved two successive inversion steps: first tosylation of 4R-OH function under Mitsunobu conditions^{10a} to yield 4S- OTs derivative



i, H₂ / Pd C ii, FmocCl / CBzCl, Na₂CO₃ iii, TFA:DCM (1:1) iv, Thymin-l-ylacetic acid, DCC HOBT v, 1M NaOH vi, Piperidine:DMF (2:8), vii, DCC HOBT

which in subsequent reaction with NaN₃ underwent an inversion to yield the azide 4R-N₃ (2b), with an overall retention of configuration at C4. This was transformed to the fully protected 4R-N₃ (2b), with an overall similar set of reactions as described for 4S, 2S isomer (3a). Paths C and D leading to 4(S/R), 2R-derivatives (3c,d) begin with an inversion step at C-2 by treatment of 1 with AcOH-Ac₂O to obtain 4R-hydroxy-2R-proline¹³, followed by an identical set of reactions as in paths A and B respectively. Thus all possible stereoisomers of 4-amino-N^{α}-thymin-1-ylacetyl prolines which are the T-monomeric units for the designed 4-aminoprolyl nucleic acids were obtained starting from 1 with overall yields in the range of 28-30%.

Although the above strategy involving condensation of prolyl ring NH with N-carboxy-methylated nucleobase was successful for synthesis of T-monomers, it is fraught with experimental difficulties for synthesis of monomers of other nucleobases, in particular that of purines.¹² We therefore sought to employ an alternative route towards a general synthesis of prolyl nucleic acid monomers which consisted of direct coupling of individual nucleobases (T, C, G and A) with suitably protected N^{α}-(bromoacetyl) proline (5). Such a strategy has been reported to be efficient in large scale synthesis of monomers of open chain peptide nucleic acids.¹⁴ In the present work, from a synthetic point of view, the 4*R*-azido-N^{α}-(bromoacetyl) proline benzyl ester 5 is convenient, since in a subsequent catalytic hydrogenation step, simultaneous transformation of 4-azido to 4-amino and deprotection of 2-benzyl ester to carboxylic acid can be achieved. It is also possible to selectively reduce 4-azido to 4-amino by catalytic transfer hydrogenation using ammonium formate/Pd-C to obtain the C-protected monomeric units. With this working rationale, we attempted the synthesis of prolyl nucleic acid monomers corresponding to A,G,C and T nucleobases by this alternative method and illustrate the feasibility of the method by a synthesis of all 4 monomers (10-13) corresponding to 4*R*,2*S* series (Scheme II).

4R-Hydroxyproline was converted into its benzyl ester by reaction with benzyl alcohol and *p*-toluenesulphonic acid¹¹ followed by protection of N^{α} with BOC¹¹ to get 4a which was transformed in two further steps to the 4*R*-azido derivative 4c. Deblocking of N^{α}-Boc with TFA afforded the free amine, which upon treatment with bromoacetylchloride in presence of aq. sodium bicarbonate, afforded the desired common intermediate 5. This reaction gave poor yields when triethylamine was employed as a base.¹⁴ The individual alkylation of the four nucleobases thymine, adenine, cytosine and 2-amino-6-chloropurine with 5 was done by reaction of the two components in DMF in presence of a suitable base, followed by purification by chromatography to obtain 6, 7, 8 and 9 respectively, in 50 to 80% yield. While the alkylation of thymine and cytosine is regiospecific, providing only N1 substitution products, that of purines is known to be non regiospecific leading to significant N7 substituted product along with the desired major product from N9-substitution.¹⁴ In the present work with adenine, no N7 regioisomer was observed as determined by NMR (see later). For synthesis of G-monomer, direct alkylation of guanine is not a facile reaction and hence 2-amino-6-chloropurine was alkylated with 5 to obtain exclusively the N9 regioisomer 9. 4*R*-Azido group in

6-9 was then hydrogenated and protected with BOC¹¹ to get the protected monomers 10-13a respectively. 13a was then hydrolysed to the desired G-monomer 13b by 1N NaOH.¹⁴ This formally completed the synthesis of



i, TFA:DCM (1:1); ii, BrCH₂COCl/ Na₂CO₃; iii, Thymine (T) and 6-Cl-2NH₂ purine/K₂CO₃, Cytosine (C) and Adenine (A)/NaH; iv, ammonium formate/Pd-C, v, BOC-N₃, Na₂CO₃, vi, 1M NaOH

all the nucleobase 4R,2S monomers required for building prolyl nucleic acids. This methodology coupled with that in Scheme-1 enables the synthesis of all possible 4-amino proline stereoisomers, corresponding to N1 (T and C) and N9 (A and G) substitution with the four nucleobases.

All compounds including those obtained as the intermediates in pathways (A-D) and in Scheme-2 were unambiguously characterized for structural purity by ¹H and ¹³C NMR. The NMR spectra of N1-acyl substituted compounds (both BOC and carboxymethylated nucleobases) showed characteristic presence of *syn* and *anti* rotamer mixtures originating from the restricted rotation about the tertiary amide bond (>N-CO). The energy barriers for such rotation are high, leading to resolution of *syn* and *anti* rotamers at ambient temperature on NMR time scale, as shown by a doubling of relevant resonances. An equilibrium composition of approximately 80:20 was noticed in most of the compounds in CDCl₃, although the assignment of specific resonances to individual rotamers was not possible in the present compounds due to complex and overlapping spectral patterns for prolyl ring protons in 200MHz spectra.

¹³C NMR is valuable in identification of N7/N9 regioisomers in alkylation reactions of purines. In case of 7- and 9- methyladenines it has been established that the N9 isomer exhibits downfield shifts of 6 and 4 ppm for C-5 and C-6 respectively and upfield shifts of 10 and 4.5 ppm for C-4 and C-6 compared to corresponding chemical shifts of N7 regioisomer.¹⁵ This pattern was also noticed with A-monomer of peptidenucleic acids¹⁴. The ¹³C NMR spectral data of compounds 7 and 11 prepared presently by alkyation of adenine was consistent with the data reported for N9 isomer. Further, the data for compound 7 prepared by two different routes (alkylation of adenine by 5 or by condensation of 4R-azido-2S proline benzyl ester with adenine-9-yl acetic acid¹²) was in complete agreement, thus supporting the assigned structures. The chiral purity of the synthesised compounds was ascertained by following optical rotation measurements at every step. The enantiomeric pairs (2a/2d)/(2b/2c) and (3a/3d)/(3b/3c) exhibited equal but opposite optical rotations, thus establishing the optical purity of compounds 2a-d and 3a-d.

Synthesis and characterization of model prolyl-nucleic acid chiral dimers 17 and 18

To examine the efficacy of synthesised monomers in polymer building reactions, the syntheses of dipeptides 17 and 18 were accomplished from 3, 3a and 3c (Scheme 1). This was done first by hydrolysis of 3 and 3c with alkali to yield the carboxylic components 14 and 16. C-4 amino deprotection of 3a with piperidine gave the common amino component 15 which was then condensed with 14 and 16 in presence of DCC-HOBT¹¹ to give the prolyl PNA dipeptides 17 (NH-4*S*,2*S*;4*S*,2*S*-OH) and 18 (NH-4*S*,2*R*;4*S*,2*S*-OH) respectively. The structural integrity of these two dipeptides was established by ¹H NMR and the diagnostic mass peaks (M+1) at 723 and 811 respectively in FAB-MS. The dipeptides 17 and 18 upon deprotection with appropriate reagents gave the free amines 17a and 18a which were characterised by ¹H NMR .

For a preliminary examination of the structural effects induced by different relationships of the chiral centres in dipeptides 17a and 18a, the CD spectra of the prolyl PNA are shown in Figure 1. These show characteristically different Cotton effects in the region 240-300 nm, with 17a having only a negative CD band at 265 nm and 18a both negative and positive CD bands at 278 and 255 nm respectively. These bands though weak, are completely absent in open chain PNA molecules⁵ and must arise in 17a and 18a due to asymmetry in nucleobase stacking induced by the chiral peptide backbone. Substantial differences are also noticed in relative CD patterns in the region 210-230 nm, originating from stereochemical differences in the peptide linkage. The two dipeptides differ in prolyl ring stereochemistry at only one centre and it is remarkable that this single change affects the sign, amplitude and band position of the CD spectrum. This is a clear evidence



Figure 1. CD spectra of 17a (_____) and 18a (----) in Tris buffer (10 mM, pH 7.2) at 25°C

for an ordered conformation of bases in dimers which senses the backbone chirality. In addition, stacking of bases could also be a function of syn/anti rotamer population due to restricted rotation around >N1-CO bond.

CONCLUSION

In this paper, we have demonstrated the design and preparation of stereochemically defined monomer building blocks, based on 4-aminoproline backbone, which are required for the synthesis of conformationally rigid, polychiral PNA. The possible effects of stereostructural constraints on backbone is preliminarily seen as induced chirality in base stacking in the model dipeptides which exhibit significant differences in their CD patterns upon change of stereochemistry even at a single site on backbone. Achiral PNA (II) has enormous structural freedom to assume a variety of conformations in solution (arising from the intervening methylene groups in the backbone) with a high statistical probability of those productive in DNA binding. The details of conformational preferences of PNA for efficient DNA binding are not yet deciphered in literature^{16,17} and the availability of PNAs with predefined rigid conformationally defined building blocks reported here along with different nucleobases (A, T, C and G) when used in appropriate rational combinations, would lead to a library of chiral PNAs with diverse backbone geometry, that are useful for hybridisation screening with appropriate DNA/RNA targets. We are currently engaged in solid phase synthesis of homooligomers from these chiral monomeric units to generate polychiral PNAs for evaluating their biophysical properties.

EXPERIMENTAL

Solvents were obtained from standard commercial sources, distilled and dried before use. Nucleobases, 4-hydroxyproline, t-butoxycarbonylchloride, Fluorenemethoxycarbonylchloride, benzyloxy carbonyl chloride, dicyclohexylcarbodiimide and hydroxybenzotriazole were obtained from Aldrich chemical company. ¹H and ¹³C NMR spectra were recorded using Bruker AC 200 MHz FTNMR spectrometer. Optical rotations were measured on JASCO DIP-181 polarimeter and CD spectra were obtained on a JASCO J600 instrument. Silica gel column chromatographic analyses were followed by pre-coated silica gel TLC plates (E. Merck 5554) and visualised by ninhydrin spray (amino compounds). All TLCs were run using DCM containing appropriate amounts of methanol.

4-Azido-N^{α}-(tert-butoxycarbonyl) proline methyl ester 2a-d: N^{α}-(tert-butoxycarbonyl-4-(*p*-toluene sulfonyloxy) proline methyl esters (4.0 g, 10 mmole) of appropriate stereochemistry (4*R*,2*S*; 4*S*,2*S*; 4*R*,2*R* and 4*S*,2*R* synthesised according to ref 10), were dissolved in dry DMF (20 mL) to which was added NaN₃ (1.2 g). The reaction mixture was stirred at 50^o C for 5 h after which the solvent was removed under reduced pressure and the residue worked up with ether. The products obtained 2a-d respectively were purified by silica gel column chromatography using pet. ether containing increasing amounts of ethyl acetate.

Compound 2a: ¹H NMR, CDCl₃, δ , 4.43 (dq, J=6.0, 8.8, 5.3 Hz, H-2), 4.33 (brm, 1H, H-4), 3.76 (s, 3H, OCH₃), 3.74 (m, 1H, H-5a), 3.48 (m, 1H, H-5b), 2.47 (m, 1H, H-3a), 2.17 (dt, J=4.3, 1.3 Hz, 1H, H-3b), ¹³C NMR, CDCl₃ δ , 172.2 (ma) and 171.9 (mi) (COOMe), 154.0 (mi) and 153.4 (ma) (OCON<), 80.5 (CMe₃), 59.3 (mi) and 58.3 (ma) (C-4), 57.7 (ma) and 57.4 (mi) (C-2), 52.3 (OMe), 51.3 (mi) and 50.8 (ma) (C-5), 36.0 (ma) and 35.1 (mi) (C-3), 28.3 [C(CH₃)₃], [α]_D²⁰ = -31.3 (c = 0.4, MeOH)

Compound **2b**, ¹H NMR, CDCl₃, δ , 4.36 (dt, J = 6.9, 7.6 Hz), 4.19 (m, 1H, H-4), 3.74 (s, 3H, OCH3), 3.64 (dd, J = 5.3, 12 Hz, H-5a), 3.47 (dt, J = 3.8, 12 Hz, H-5b), 2.32 (m, 1H, H-3a), 2.16 (m, 1H, H-3b), 1.46 (ma) and 1.41 (mi) [s, 9H, C(CH)₃]. ¹³C NMR, CDCl₃, δ 172.5 (ma) and 172.2 (mi) (CO), 153.6 (mi) and 152.9 (ma) (NCO), 80 (CCH₃)₃, 59.04 (mi) and 58.5 (ma) (C-4), 51.7 (OCH₃), 51.1 (mi) and 50.9 (ma), C-5, 35.9 (ma) and 35.0 (mi) (C-3), 27.8 (C-CMe₃), $[\alpha]_D^{20}$ -60.9 (c = 0.4, MeOH)

Compound 2c, ¹H NMR and ¹³C NMR same as 2b, $[\alpha]_D^{20}$ +61.2 (c = 0.4, MeOH)

Compound 2d, ¹H NMR and ¹³C NMR same as 2a, $[\alpha]_D^{20}$ +32.0 (c = 0.4, MeOH)

4N-(Fluorenylmethoxycarbonyl)-N^{α}-(thymin-1-ylacetyl) proline methyl ester 3a-d: The compounds 2a-d of stereochemistry 4*S*,2*S*; 4*R*,2*S*; 4*R*,2*R* and 4*R*,2*R* respectively were converted to 3a-d by the following set of reactions: (i) hydrogenation of substrate (1.4g) over Pd-C (0.15 g) at 30 psi in MeOH for 4 h to yield 4-aminoproline derivatives, (ii) conversion of these to corresponding 4-N-(fluorenylmethoxycarbonyl)-N-(t-

benzyloxycarbonyl chloride (0.34 g, 2 mmole) in a mixture of dioxane (3 mL) and 10% aq. sodium carbonate (5 mL) and stirring at ambient temperature for 6 h. (iii) deprotection of N-BOC with 50% TFA:DCM¹¹ (1 h, 25° C) and (iv) condensation of resulting 4-N(fluorenylmethoxycarbonyl) proline methyl esters (2.0 g, 5 mmole) with thymine-1-ylacetic acid (1.1 g, 6 mmole) in dry DMF:DCM (1:1, 40 mL) in presence of DCC (1.6 g, 6.5 mmole) and HOBT (0.75 g, 6 mmole) at 4°C for 24 h. The precipitated DCU was filtered off, washed with DCM and the residue after concentration was worked up in DCM (2x50 mL) by successively washing with aq. NaHCO₃ (2x20 mL), aq. KHSO₄ (20 mL), brine (20 mL) and water (20 mL). The dried organic extract upon concentration and column purification (elution DCM:MeOH) gave the products **3a-d** in 30% overall yield.

Compound 3: ¹H NMR, CDCl₃ δ 9.02 (mi) & 8.95 (ma) (s, 1H,*T* NH), 7.4 (s, 5H, Ar-H), 7.09 (mi) & 7.0 (ma) (s, 1H, *T* H-6), 6.0 (d, 1H, C-4 NH), 5.1 (s, 2H, ArCH₂), 4.65 (d, 1H, J = 16 Hz, *T* CHa) 4.5-4.44 (m, 2H, H-2, H-4), 4.25 (d, 1H, J = 16 Hz, *T* CHb), 4.0-3.5 (m, 2H, H-5), 3.75 (ma), 3.7 (mi) (s, 3H, OCH₃), 2.5 (ma) & 2.35 (mi) (m, 1H, H-3b), 2.05 (mi) & 2.0 (ma) (br, 1H, H-3a), 1.9 (s, 3H, *T* CH₃) ¹³C NMR, CDCl₃ δ 155.8 (*T* C-2), 151.4 (*T* C-4), 141.2 (*T* C-6), 110.5 (*T* C-5), 58.0 (OCH₃), 52.67 (C-2), 48.7 (*T* CH₂), 50.9 (C-4), 43.6 (C-5), 35.0 (C-3), 12.2 (*T* CH₃).

Compound 3a: ¹H NMR, CDCl₃ δ , 8.62 (s, 1H, *T* NH), 7.76-7.26 (m, 8H, Fmoc), 7.07 (mi) & 7.05 (ma) (s, 1H, *T* H-6) 5.98 (ma) (d, J = 11 Hz, C-4 NH) & 5.3 (mi, br, C-4 NH), 4.73-4.16 (m, 7H, H-2, H-4, *T* CH₂, Fmoc CH and OCH₂), 4.0-3.57 (br-m, 2H, H-5), 3.78 (ma) & 3.76 (mi) (s, 3H, OCH₃), 2.5 (br-m, 1H, H-3b), 2.04 (m, 1H, H-3a) & 1.93 (s, 3H, *T* CH₃). ¹³C NMR, CDCl₃ δ , 155.8 (*T* C-2), 151.3 (*T* C-4), 141.2 (*T* C-6), 110.4 (*T* C-5), 57.9 (OCH₃), 52.6 (C-2), 51.9 (*T* CH₂), 50.8 (C-4), 48.6 (C-5), 34.5 (C-3), 12.2 (*T* CH₃). [α]_D²⁰ (c = 0.4, MeOH) = -38.8°, FAB-MS:[M+1] = 533.

Compound **3b**: ¹H NMR CDCl₃ δ ; 9.75 (s, 1H, T NH), 7.76-7.25 (m, 8H, Fmoc), 7.02 (mi) & 6.96 (ma) (s, 1H, *T* H-6), 6.34 (ma) & 6.12 (mi) (d, 1H, J = 7.4 Hz, C-4 NH), 4.82-3.89 (m, 7H, H-2, H-4, *T* CH₂, Fmoc CH, OCH₂), 3.72 (s, 3H, OCH₃), 3.8-3.66 (m, 2H, H-5), 2.44 (mi) & 2.17 (ma) (br-m, 2H, H-3a,b), 2.1 (s, 3H, *T* CH₃). ¹³C NMR CDCl₃ δ , 156.1 (*T* C-2), 151.6 (*T* C-4), 141.3 (*T* C-6), 110.8 (*T* C-5), 57.9 (OCH₃), 52.6 (C-2), 51.4 (*T* CH₂), 50.7 (C-4), 48.9 (C-5), 35.0 (C-3), 12.2 (*T* CH₃), $[\alpha]_D^{20}$ (c = 0.4, MeOH) = +22.5°, FAB-MS: [M+1] = 533.

Compound 3c: ¹H NMR (CDCl₃) δ : 9.82 (s, 1H, T NH), 7.76-7.25 (m, 8H, Fmoc), 7.02 (mi) & 6.94 (ma) (s, 1H, T H-6), 6.4 (ma) & 6.18 (mi) (d, J = 7.4 Hz, C-4 NH), 4.83-3.89 (m, 7H, H-2, H-4, T CH₂, Fmoc CH, CH₂), 3.71 (ma) & 3.73 (mi) (s, 3H, OCH₃), 3.78-3.72 (m, 2H, H-5), 2.44 (mi) & 2.22 (ma) (brm, 2H, H-3a,b), 2.11 (s, 3H, T CH₃). ¹³C NMR (CDCl₃) δ : 156.4 T C-2), 151.9 (T C-4), 141.5 (T C-6), 111.3 (T C-5),

58.2 (OCH₃), 52.9 (C-2), 51.9 (*T* CH₂), 51.0 (C-4), 49.2 (C-5), 35.4 (C-3), 12.6 (*T* CH₃). $[\alpha]_D^{20} = -22.8$ (c = 0.4, MeOH), FAB-MS: [M+1] = 533.

Compound **3d:** ¹H NMR, CDCl₃ δ : 7.76-7.26 (m, 8H, Fmoc), 7.08 (mi) & 7.0 (ma) (s, 1H, *T* H-6), 6.05 (ma) & 5.4 (mi) (d, 2H, J = 7.4 Hz, C-4 NH), 4.75-4.15 (m, 7H, H-2, H-4, *T* CH₂, Fmoc CH, CH₂), 3.98-3.56 (br-m, 2H, H-5) 3.77 (ma) & 3.75 (mi) (s, 3H, OCH₃), 2.5 (br-m, 2H, H-3a,b), 1.93 (s, 3H, *T* CH₃). ¹³C NMR CDCl₃ δ : 156.0 (*T* C-2), 151.6 (*T* C-4), 141.4 (*T* C-6), 110.6 (*T*-C5), 58.1 (OCH₃), 52.8 (C-2), 52.0 (*T* CH₂), 51.0(C-4), 48.9 (C-5), 34.8 C-3), 12.3 (*T* CH₃). $[\alpha]_D^{20} = +38.63$, (c = 0.4, MeOH), FAB-MS: [M+1]=533.

4S-(p-Toluenesulfonyl)-N^{\alpha}-(tertbutoxycarbonyl) proline benzyl ester 4b: N-(t-Butoxycarbonyl)-4*R*hydroxy-2*S* proline benzylester¹¹ 4a was converted to corresponding 4*S*-tosyl derivative 4b under Mitsunobu conditions using triphenyl phosphine (9.2 g), DEAD (6.2 g) in THF (300 mL) and methyl-*p*-toluenesulphonate (6.6 g) in THF (60 mL) The product was purified by column chromatography, eluting with pet ether-EtOAc to obtain 4b in 60% yield. ¹H NMR, CDCl₃, δ , 7.25-7.85 (q, 9H, overlapping signals ArH), 5.3 (m, 3H, overlapping signals for ArCH₂ and H-4), 4.45 (m, 1H, H-2), 3.65 (m, 2H, H-5a,b), 2.45 (s, 3H, ArCH₃), 2.3-2.55 (m, 2H, H-3), 1.38 and 1.28 [2 x s, 9H, C(CH₃)₃]

4*R*-Azido-N^{α}-(tertbutoxycarbonyl)-2*S*-proline benzyl ester 4c: Compound 4b (4.6 g, 10 mmole), was dissolved in DMF (25 mL) containing NaN₃ (1.2 g, 20 mmole) and stirred at 50⁰ C for 3 h. Work up and column chromatography afforded 4c (2.8 g, 80%) ¹H NMR, (CDCl₃), δ , 7.35 (s, 5H, ArH), 5.15 (m, 2H, ArCH₂), 4.45 (mi) and 4.34 (ma) (2 x t, 1H, J = 7.5 Hz, H-2), 4.13 (brm, 1H, H-4), 3.66 (m, 1H, H-5a), 3.56 (ma, dd, J=2.8, 11.4 Hz, H-5b), 3.42 (mi, dd, J=3.5, 11.4 Hz, H-5a), 3.56 (ma, dd, J=2.8, 11.4 Hz, H-5b), 2.3 (m, 1H, H-3a), 2.13 (m, 1H, H-3b).

4*R***-Azido-N^{\alpha}-(bromoacetyl) proline benzyl ester 5:** Compound 4c (3.5 g) was treated with 50% TFA in DCM for 1 h and the residue obtained after evaporation of solvent was dissolved in dioxane (15 mL) containing 10% aq. Na₂CO₃ (25 mL, pH 8.0). The mixture was cooled in ice bath for addition of bromoacetyl chloride (7 equiv). The pH was then adjusted to 8.0 by addition of aq. Na₂CO₃. The mixture was concentrated to half its volume and extracted with DCM. The dried organic layer upon concentration and column purification gave 5 (3.1 g, 80% yield) ¹H NMR, (CDCl₃) (2 rotamers), δ , 7.37 (s, 5H, ArH), 5.16 (m, 2H, ArCH₂), 4.72 (mi) and 4.63 (ma) (2xt, 1H, J = 7.0 Hz, H-2), 4.3 (m, 1H, H-4), 3.92 (m, 1H, H-5a), 3.65 (m, 1H, H-5b), 3.81 (s, 2H, CH₂Br), 2.36 (m, 1H, H-3a), 2.18 (m, 1H, H-3b)

4*R*-Azido-N^{α}-(thymin-1-ylacetyl) proline benzyl ester 6: A mixture of 5 (1.25 g, 0.3 mmole), thymine (0.6 g, 0.3 mmole) and solid K₂CO₃ (0.7 g, 3 mmole) in dry DMF (20 mL) was stirred at 25°C for 3 h. The residue obtained after aqueous work up was purified by column chromatography by elution with DCM-MeOH to obtain 6 (yield 90%), ¹H NMR, (CDCl₃) δ , 9.45 (brs, 1H, NH), 7.38 (mi) and 7.32 (ma) (s, 5H, Ar-H), 7.05 (ma) and 6.85 (mi) (s, 1H, T-CH), 5.2 (m, 2H, ArCH₂), 4.84 (mi, q, J = 5.8, 7.8 Hz, H-2), 4.65 (ma, q, J = 13.7, 2.0 Hz, H-2), 4.6 (d, 1H, J=16.2, T C-Ha), 4.37 (d, 1H, J=16.2 Hz, T C-Hb), 4.35 (ma) and 4.2 (mi) (m,

1H, H-4), 3.92 (dd, 1H, J = 11.6, 5.8 Hz, H-5a), 3.7 (dt, J = 11, 4.0 Hz, 1H, H-5b), 2.35 (m, 2H, H-3a.b), 1.95 (s, 3H, T CH₃).

4R-Azido-N^{α}-(adenin-9-ylacetyl) proline benzyl ester 7, 4R-azido-N^{α}-(cytosin-1-ylacetyl) proline benzyl ester 8 and 4R-Azido-N^{α}-(6chloro-2-aminopurin-9-ylacetyl) proline benzyl ester 9: These were synthesised from 5, appropriate nucleobase and either NaH (for C and A) or K₂CO₃ (for 2-amino-6-chloropurine) by following a similar procedure as above. The products 7, 8 and 9 were purified by column chromatography by elution with DCM-MeOH (yields 50-70%).

Compound 7, ¹H NMR,(CDCl₃) δ 8.31 (ma) and 8.3 (mi) (s, 1H, A H-8), 7.93 (ma) and 7.81 (mi) (s, 1H, A H-2), 7.38 (mi) and 7.33 (ma) (s, 5H, ArH), 6.36 (ma) and 6.26 (mi) (brs, Ad NH₂), 5.25 (mi) and 5.15 (ma) (2 x dd, J = 12.5 Hz, 2H, ArCH₂), 4.95 (ma, s, A CH₂) and 4.92 (mi, t, J = 7.5 Hz, A CH₂), 4.65 (t, 1H, J = 9 Hz, H-2), 4.35 (ma) and 4.18 (mi) (m, 1H, H-4), 3.92 (q, 1H, J = 5.2, 10 Hz, H-5a), 3.71 (q, 1H, J=2.5, 10 Hz, H-5b), 2.35 (m, 2H, H-3a,b) ¹³C NMR, (CDCl₃) δ , 170.6 (COOBn), 165.2 (mi) and 164.7 (ma) (>NCO), 155.5 (A C-6), 152.7 (A C-2), 149.7 (A C-4), 141.1 (A C-8), 134.9-127.9 (Ar), 118.5 (A C-5), 68.02 (mi) and 67.1 (ma) (ArCH₂), 59.3 (C-2), 57.9 (ma) and 57.2 (mi) (C-4), 51.3 (A CH₂), 44.3 (ma) and 43.8 (mi) (C-5), 36.4 (mi) and 34.08 (ma) (C-3), [α]_D²⁰ -44.0 (c = 0.3, MeOH)

Compound 8, ¹H NMR, (CDCl₃+D₂O), δ 7.45-7.15 (m, 6H, ArH and *C* H-6), 5.9 (d, J = 8.0 Hz, *C* H-5), 4.95-5.53 (m, 2H, ArCH₂), 4.82-4.45 (overlapping multiplets, 3H, N-CH₂, H-2), 4.35 (brm, 1H, H-4), 3.95 (brm, 1H, H-5b), 3.7 (brm, 1H, H-5a), 2.05-2.55 (overlapping multiplet, 2H, H-3a,b), $[\alpha]_D^{20}$ -51.3 (c = 0.3, MeOH) *Compound* 9, ¹H NMR, (CDCl₃), δ 8.0 (mi) and 7.8 (ma) (s, H-8), 5.31 (brs, 2H, NH₂), 5.23 (mi, m, ArCH₂), 5.11 (dd, J=12.2, ArCH₂), 4.6 (mi, m, NCH₂) and 4.61 (ma, t, J = 7.6 Hz, NCH₂), 4.35 (ma, m, H-4) and 4.2 (mi, m, H-4), 3.89 (ma, dd, J = 5.4, 10.2 Hz, H-5a) and 3.81 (mi, dd, J = 3.1, 12.9 Hz, H-5a), 3.64 (dd, J = 3.3, 10.1 Hz, H-5b), 2.4 (mi, m, H-3a), 2.37 (ma, dq, J = 14.0, 7.5 Hz, H-3a), 2.17 (dq, J = 14, 7.5 Hz, H-3b), $[\alpha]_D^{20}$ -42.5 (c = 0.4, MeOH).

4*R*-N-(ter/butoxycarbonyl)-N^{α}-(thymin-1ylacetyl) proline, 10: Compound 6 (2 g) in MeOH was hydrogenated under pressure (30 psi) using Pd-C (10-12%, 200 mg) as catalyst for 15 h. The catalyst was filtered and the free amino acid product 10, was isolated by evaporation of solvent. This compound (1.4 g, 5 mmole) was dissolved in dioxane:water (1:1, 20 mL) and treated with tertbutoxycarbonyl azide (0.8 mL) and kept stirred at 25° C for 24 h. with pH maintained at 9.0 by addition of 4N NaOH. The reaction mixture was concentrated to 10 mL, neutralised with Dowex 50 H⁺ and filtered. Removal of solvent from the filtrate afforded the required product which was purified by crystalisation from MeOH. ¹H NMR, (D₂O), δ 7.39 (d, J = 1.2 Hz, *T* H-6), 4.7 (t, J = 12.1 Hz, N-CH₂), 4.52 (t, J = 9.2 Hz, H-2), 4.36 (m, 1H, H-4), 4.2 (mi) and 3.95 (ma) (dd, H-5b), 2.25-2.5 (bt, 2H, H-3a,b), 1.9 (s, 3H, *T* CH3), 1.46 [s, 9H, C(CH₃)₃], [α]_D²⁰ -14.7 (c = 0.3, MeOH).

4R-N-(tertbutoxycarbonyl)-N^{α}-(A/G/C-ylacetyl) proline benzyl ester 11-13a: The 4R-azido compounds 7-9 (1.0 g) were individually dissolved in methanol (50 m L) and reacted with ammonium formate (4 eq) and PdC (10-12%) for 24 h. after which the catalyst was removed by filtration over celite. The filtrate was evaporated under vacuo and the residue dissolved in water: dioxane (1:1, 15 mL), was treated with 1-butoxycarbonylazide (1.5 eq) and TEA (1.5 eq) at 50° C for 15 h, after which the products 11-13a, were isolated by chromatographic purification (yields 60-70%).

Compound 11, ¹H NMR,(CDCl₃), δ 8.29 (s, 1H, *A* H-8), 7.82 (ma) and 7.72 (mi) (s, 1H, H-2), 7.32-7.36 (m, 5H, ArH, 6.48 (brs, 2H, NH₂), 5.75 (brd, ma) and 5.52 (br, mi, 1H, NHBOC), 5.25 (mi) and 5.13 (ma) (dd, 2H, ArCH₂), 4.9 (dd, J = 15.6 Hz, 2H, Ad CH₂), 4.64 (t, 1H, J = 5.8 Hz, H-2), 4.36 (br, 1H, C-4), 3.96 (dd, 1H, H-5a), 3.78 (brm, 1H, H-5b), 2.5-2.05 (brm, 2H, H-3a,b), 1.49 [s, 9H, C(CH₃)₃], ¹³C NMR, (CDCl₃), δ 171.4 (ma) and 171.1 (mi) (COOR), 165.5 (CONH), 155.4 (C6 and OCON<), 152.4 (Ad C-2), 149.3 (*A* C-4), 141.6 (*A* C-8), 135.1-127 (Ar), 118.2 (*A* C-5), 79.8 (CMe₃), 67.9 (mi) and 67.1(ma) (ArCH₂), 57.8 (C-2), 51.4 (C-4), 50.0 (*A* NCH₂), 44.7 (C-5), 34.8 (C-3), 28.3 [C(CH₃)₃], [α]_D²⁰ -20.7 (c = 0.3, MeOH).

Compound 12. ¹H NMR, (CDCl₃+D₂O), δ 7.25-7.4 (overlapping signals, ArH, C H-6), 5.9 (d, J = 6.36 Hz, 1H, C H-5), 5.1 (overlapping q, 2H, ArCH₂), 4.55-4.95 (brm, 2H, N-CH₂), 4.15-4.5 (brm, 2H, H-3a,b), 1.45 [s, 9H, C(CH₃)₃], [α]_D²⁰ -49.6 (c = 0.3, MeOH).

Compound 13a, ¹H NMR, (CDCl₃), δ 7.85 (ma) and 7.7 (mi) (s, 1H, G H-8), 7.3 (m, 5H, ArH), 4.97-5.45 (overlapping signals, 4H, NH₂ and ArCH₂), 4.82 (dd, 2H, N-CH₂), 4.65 t, J = 4.2 Hz, H-2), 4.4 (brm, 1H, H-4), 4.05 (brm, 1H, H-5a), 3.65 (brm, 1H, H-5b), 2.45 (mi, m) and 2.22 (ma, t) (2H, H-3a,b), 1.47 [s, 9H, C(CH₃)₃], [α]_D²⁰ -20.5 (c = 0.2, MeOH).

4*R*-N-(tertbutoxycarbonyl)-N^{α}- (guanin-9-ylacetyl) proline, 13b: A mixture of 13a (0.5g, 1mmole) and 1N aq. NaOH (2mL) was stirred at r.t for 2 h after which it was neutralised with ion-exchange resin, Dowex H⁺. MeOH (5 mL) was added, the resin was filtered off and the product that slowly precipitated was collected (yield 60%).¹H NMR, (*d*₆-DMSO:D₂O), δ 7.7 (s, 1H, *G* H-8), 4.92 (q, 2H, J = 19.5 Hz, N-CH₂), 4.28 (q, 1H, J = 8.1, 4.8 Hz, H-2), 4.18 (t, 1H, J = 7.2 Hz, H-4), 3.84 (q, 1H, J = 9.6, 5.8 Hz, H-5a), 3.4 (q, 1H, J = 9.6, 7.2, H-5b), 2.1 (dq, 1H, H-3a,3b), 1.4 [s, 9H, C(CH₃)₃]

Dipeptide Nucleic acids 17a and 18a: General method: The thymine monomers 14 and 16 (carboxyl component, 0.12 mmole) were condensed separately with the amino components 15 (30 mg, 0.1 mmole) in DMF (500 μ L) in presence of DCC (25 mg, 0.12 mmole) and HOBT (15 mg, 0.1 mmole) at 25° C for 4 h. The usual work up afforded the protected dipeptide products 17 and 18 respectively, which were quantitatively deprotected in one step using either H₂/Pd-C or piperidine:DMF to yield the free dipeptides 17a and 18a respectively.

Compound 17a: ¹H NMR (500 MHz, D₂O) δ : 7.38 (2H, T CH), 4.7-4.28 (8H, T CH₂ x 2, H-4, H-2 x 2), 4.16-3.67 (m, 4H, H-5a,b x 2), 3.75 (s, 3H, OCH₃), 2.8-2.12 (m, 4H, H-3a,b x 2), 1.88 (s, 6H, T CH₃ x 2). UV $\lambda_{max} = 269$ nm. *Compound* 18a: ¹H NMR (500 MHz, D₂O) δ: 7.42,7.38 (2H, *T* CH), 4.8-4.4 (8H, *T* CH₂ x 2, H-4, H-2 x2), 4.2-3.7 (m, 4H, H-5a,b x 2), 3.77, 3.73 (s, 3H, OCH₃), 2.8-2.12 (m, 4H, H-3a,b x 2), 1.89,1.87 (s, 6H, *T* CH₃ x 2). UV λ_{max}=270 nm.

Compound 17 : ¹H NMR d₆-DMSO δ : 11.3 (s, 2H, *T* NH x 2, 5.03 (s, 2H, CBz CH₂), 4.57-3.81 (m, 8H, H-2, H-4 x 2, *T* CH₂ X 2), 3.56m (s, 3H, OCH₃), 3.4-3.2 (br, 4H, H-5a,b x 2 HO), 2.3-2.55 (br, 4H, H-3a,b, x 2) 1.75 (s, 6H, *T* CH₃ x 2) FAB-MS: [M+1] = 723.

Compound 18: ¹H NMR *d*₆-acetone δ: 7.95-7.25 (m, 10H, Fmoc-Ar, *T* CH), 4.67-4.16 (m, 10H, H-2, H-4 x 2, Fmoc CH, CH₂, *T* CH₂ x 2), 3.91-3.54 (m, 4H, H-5a,b x 2), 3.7(s, 3H, OCH₃), 2.65-2.2 (br, 4H, H-3a,b x 2), 1.82, 1.8 (s,6H, *T* CH₃ x2) FAB-MS: [M+1] = 811

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