

gel electrophoresis (SDS-PAGE); the gels showed no major impurities, only a multitude of very minor impurity bands. Routine assays were conducted by monitoring tritium wash-out from [2,3-³H]- γ -butyrobetaine by a modification of the method of England et al.³⁴ Assay mixtures (pH 7.0) contained an aliquot of enzyme solution, 0.5 μ mol of [2,3-³H]- γ -butyrobetaine, 0.75 μ mol of α -ketoglutarate, 4.3 μ mol of ascorbate, 0.5 μ mol of ferrous sulfate, 1 mg of catalase, and 12.5 μ mol of potassium phosphate in a final volume of 0.25 mL. All assay components except the radiolabeled substrate were mixed in open test tubes (13 \times 100 mm) and preincubated for 3 min at 37 $^{\circ}$ C with rotary shaking (160 rpm). The substrate was added, and the incubation was continued for 45 min. The reaction was terminated by addition of 2,2'-bipyridyl (50 μ L of a 0.5 M ethanolic solution). The mixture was applied to a small column (1-mL total volume) of Dowex 50X8-400 (H⁺), and the column was eluted with 1.5 mL of water. Aliquots of the effluent were then analyzed for tritium radioactivity. In determination of the enzyme activity, two correction factors must be applied to account for the fact that (a) only one-fourth of the tritium in the labeled substrate is in the 3R position, which is the site of hydroxylation, and (b) there is a small primary tritium kinetic isotope effect for the hydroxylation, $T(V/K) = 1.5$.¹⁰ Under these conditions our most active preparations of BB hydroxylase showed a specific activity of 6.1 μ mol min⁻¹ mg⁻¹. More typically, the specific activity was about 2 μ mol min⁻¹ mg⁻¹, but there was no difference in the purity of the various preparations as judged by SDS-PAGE.

Reversible Inhibition Studies. For the determination of reversible inhibition of BB hydroxylase by substrate analogues, a series of routine assays were performed that contained various concentrations of [2,3-³H]- γ -butyrobetaine and various concentrations of the potential inhibitor.

For these assays, all components except the substrate and inhibitor were mixed and preincubated as described above, and then the inhibitor was added, followed immediately by the substrate to initiate the reaction. In this way the inhibitors could have no effect on the activation of the enzyme during the preincubation period. The remainder of the assay procedures were as above, and the inhibition data were analyzed in the form of Lineweaver-Burk plots.¹⁹

Time-Dependent Inactivation Studies. For the determination of time-dependent inactivation of BB dioxygenase by substrate analogues, the following procedure was used. A series of reaction mixtures were prepared, each of which contained (at pH 7.0) 100 μ g of BB dioxygenase, 1.2 μ mol of α -ketoglutarate, 4.5 μ mol of ascorbate, 0.6 μ mol of ferrous sulfate, 1 mg of catalase, and 50 μ mol of potassium phosphate in a final volume of 0.28 mL. After preincubation for 3 min at 37 $^{\circ}$ C with rotary shaking (160 rpm), 20- μ L aliquots of inhibitor stock solutions of appropriate concentrations were added. The incubations were continued, and at 10-min intervals aliquots were taken from the mixtures for the assay of remaining enzyme activity by using the standard routine assay procedure.

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Registry No. [³H]-1, 124201-55-4; 3, 24499-80-7; [²H]-3, 124201-56-5; [³H]-3, 124201-57-6; 4, 124201-52-1; 5, 124201-53-2; 6, 124201-54-3; D₂, 7782-39-0; T₂, 10028-17-8; γ -amino[2,3-³H]butyric acid, 13048-68-5; diethyl malonate, 105-53-3; 3,3-dimethylbutanal, 2987-16-8; methyl 5-methyl-4-hexenoate, 35901-76-9; BB hydroxylase, 9045-31-2.

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Man-Designed Bleomycin with Altered Sequence Specificity in DNA Cleavage[†]

Masami Otsuka,[‡] Takeshi Masuda,[‡] Andreas Haupt,^{‡,§} Masaji Ohno,^{*,‡} Takashi Shiraki,^{||} Yukio Sugiura,^{||} and Kenji Maeda[‡]

Contribution from the Faculty of Pharmaceutical Sciences, University of Tokyo, Hongo, Bunkyo-ku, Tokyo 113, Japan, Institute for Chemical Research, Kyoto University, Uji, Kyoto 611, Japan, and Institute of Microbial Chemistry, Kamiosaki, Shinagawa-ku, Tokyo 141, Japan. Received April 25, 1989

Abstract: The synthetic approach to the concerted antitumor mechanism of bleomycin is studied by introducing a dynamic change into the O₂-activation moiety and DNA-binding site. A model PYML(6)-bleomycin previously reported, possessing an oxygen-activating methoxy-pyridine moiety and a DNA-binding bithiazole moiety, exhibits a nucleotide cleavage mode virtually identical with that of bleomycin. Herein reported is a newly designed bleomycin analogue, PYML(6)-(4R-APA)-distamycin, wherein the 4-methoxy-pyridine moiety and a DNA-binding distamycin component are connected through an (R)-4-aminopentanoic acid linker moiety. Synthesis of PYML(6)-(4R-APA)-distamycin is carried out by condensation of the hydroxyhistidine-pentanoic acid fragment with the methoxy-pyridine moiety, followed by introduction of the distamycin moiety. PYML(6)-(4R-APA)-distamycin cleaves a G4 phage DNA fragment (100 base pairs) at 1 μ M concentration in the presence of Fe(II), oxygen, and dithiothreitol and induces dramatically altered adenine/thymine specificity. It is indicated that the specific recognition of base sequences for the cleavage is mainly controlled by the DNA affinity site and that the (R)-4-aminopentanoic acid linker seems to determine the proper arrangement of the iron-oxygen site and the distamycin moiety on DNA.

Bleomycins (BLMs) are chemotherapeutic agents used for the clinical treatment of Hodgkin's lymphoma, carcinomas of the skin, head, and neck, and tumors of the testis.¹ The drug was isolated from *Streptomyces verticillus* as a copper chelate by Umezawa

and his co-workers in 1966, and the structure was shown to be a glycopeptide consisting of an unusual hexapeptide and a disaccharide.² In addition to its clinical usefulness, BLM attracts current interest because of its unique biochemical functions. It has been well documented that BLM cleaves DNA preferentially at G-C (5' \rightarrow 3') and G-T (5' \rightarrow 3') sequences in the presence of oxygen and ferrous ion.³ The biochemical capability of BLM

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[‡] University of Tokyo.

[§] Postdoctoral fellow of Japan Society for the Promotion of Science, 1987-1988. Present address: BASF AG, Ludwigshafen, Federal Republic of Germany.

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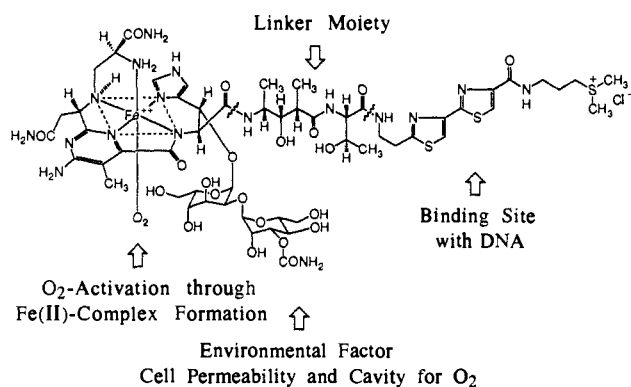


Figure 1. Proposed structure of BLM-Fe(II)-O₂ and assumed role of each functional moiety.

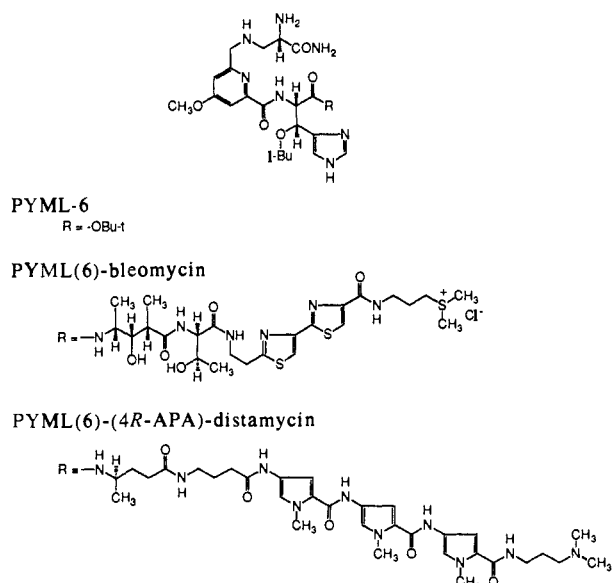


Figure 2. Synthetic analogues of BLM.

appears to be associated with the functions of the following structural units: (i) an iron-binding site (amine-pyrimidine-imidazole region) to activate dioxygen, (ii) a disaccharide moiety to facilitate the iron-oxygen complex formation and membrane transport, (iii) a DNA affinity site (bithiazole-terminal amine region) to interact with certain sequences of nucleotides, and (iv) a linker moiety to combine each part of BLM at an appropriate distance and in a good orientation (Figure 1).

In our continuing study toward man-designed BLMs, we demonstrated that the 4-aminopyrimidine nucleus and the disaccharide of BLM can be replaced by a 4-methoxypyridine and a *tert*-butyl group, respectively, and that such a synthetic model, PYML(6), was able to activate molecular oxygen as efficiently as BLM (Figure 2).⁴ Furthermore, a BLM-type DNA affinity site was coupled with PYML(6) to afford a DNA-cleaving molecule, PYML(6)-bleomycin, which showed a nucleotide sequence cleavage mode virtually identical with that of BLM *in vitro* (Figure 2).⁵ Since PYML(6) showed no significant DNA-cleaving activity, the bithiazole moiety seemed to contribute mainly to the binding to DNA by recognizing the base sequence, whereas

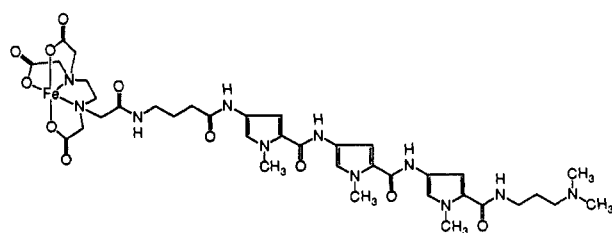
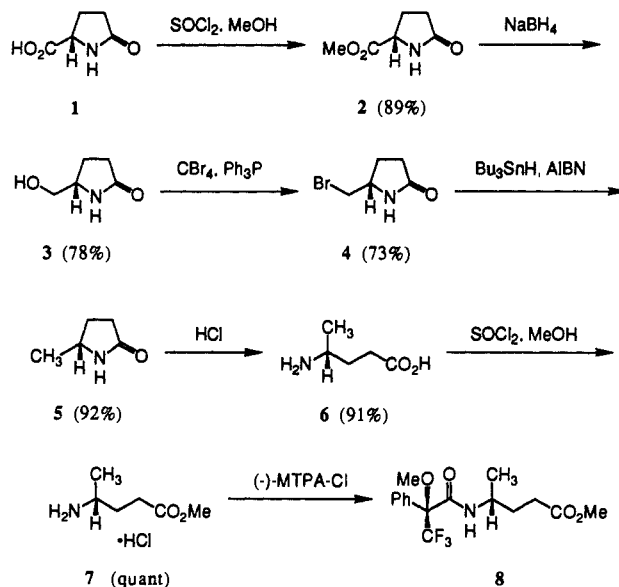


Figure 3. Distamycin-EDTA-Fe(II).

Scheme I



the sugar moiety is not directly concerned with recognition of the base sequence. In fact, a DNA-cleavage inhibition analysis and computer model-building study indicated that metallo-BLM binds in the minor groove of B-DNA, probably by the formation of hydrogen bonds between the nitrogen atoms in the bithiazole moiety and the 2-amino group of the guanine base.⁶ On the other hand, a DNA-cleaving molecule of entirely different structure, hemin-intercalator, reported by Shudo *et al.*, also exhibited a DNA-cleaving pattern remarkably similar to that of BLM.⁷ Thus it remains an alternate possibility that certain nucleotide residues are intrinsically vulnerable and sensitive to the activated oxygens generated by such iron complexes. In order to clarify whether the sequence specificity is actually dependent upon the DNA-binding site or not, we studied the relationships between the DNA-cleavage pattern and the DNA-binding moiety by modifying our synthetic model.

First, we considered the possibility that the G specificity of PYML(6)-bleomycin could be altered by replacing its bithiazole moiety with repeating *N*-methylpyrrole subunits, as envisaged in the structures of distamycin or netropsin, whose binding to A/T-rich nucleotide sequences has been unequivocally demonstrated by X-ray studies.⁸ The base specificity of distamycin or netropsin was explained by van der Waals interaction between the CH group of the *N*-methylpyrrole ring and the adenine C2 hydrogens.⁸ Furthermore, distamycin serves as a potential DNA affinity site with AT specificity, as seen in the ingenious design of distamycin-EDTA-Fe(II) by Dervan (Figure 3).⁹ Distamycin-EDTA-Fe(II) reagent was shown to cleave double-helical nucleotide sequences adjacent to 5-base-pair AT recognition sites, and the

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cleavage takes place at contiguous multiple bases, presumably because of the diffusible nature of the oxidizing species.

On the other hand, natural BLM has (2*S*,3*S*,4*R*)-4-amino-3-hydroxy-2-methylpentanoic acid as a linker between the metal-binding site and the bithiazole moiety. Although the exact role of the linker moiety remains to be clarified, it is considered that the preferred conformation of this moiety may contribute to making the flexible BLM molecule rather inflexible in the twisted minor groove of DNA. Umezawa et al. extensively studied modification of the linker moiety and clearly demonstrated that a modified BLM possessing a simplified (*R*)-4-aminopentanoic acid linker [4*R*-APA (6)]; see Scheme I] shows outstanding DNA-cleaving activity comparable to that of natural BLM.¹⁰ Their results indicated that the 4*R* configuration is particularly important among the three consecutive 2*S*,3*S*,4*R* asymmetric centers in the linear amino acid.

Therefore, we designed PYML(6)-(4*R*-APA)-distamycin, which has a PYML(6) metal core, a distamycin moiety, and a 4*R*-APA linker (Figure 2).

Results

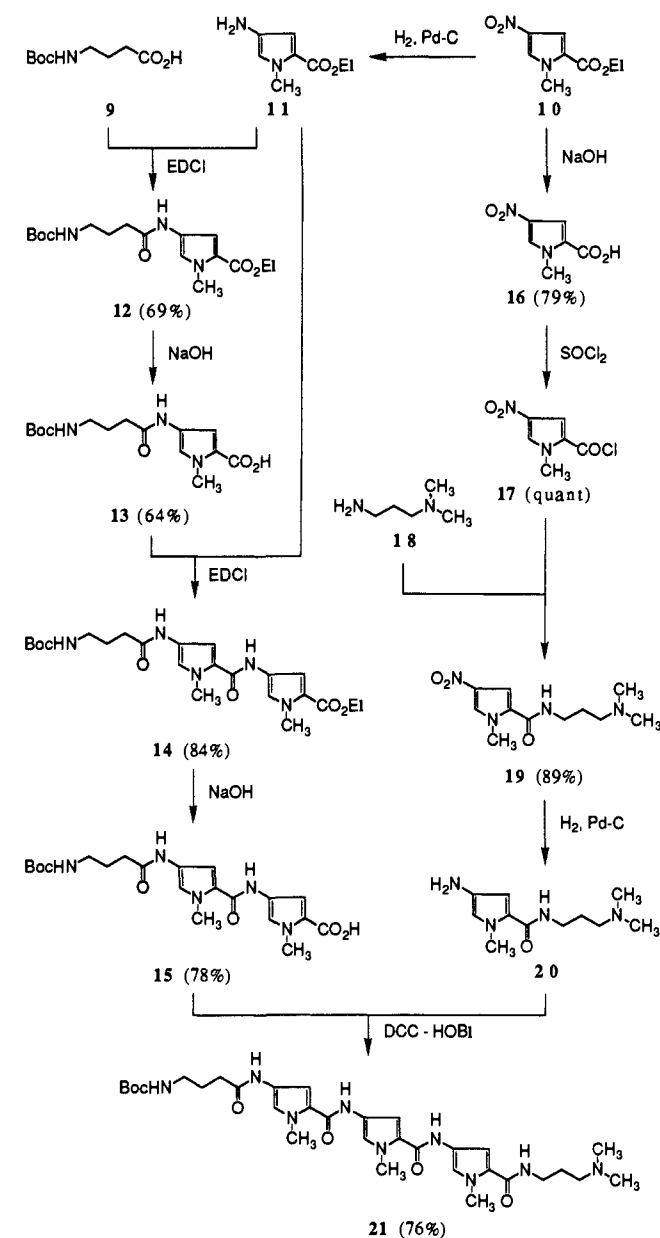
Synthesis of PYML(6)-(4*R*-APA)-Distamycin. The 4-methoxypyridine moiety and *erythro*- β -hydroxy-L-histidine part were previously prepared in the synthesis of PYML(6) and PYML(6)-bleomycin.^{4,5}

4*R*-APA was previously synthesized by Saito by Wittig reaction of an aldehyde derived from D-alanine.¹¹ We prepared 4*R*-APA (6) by a different and practical route involving a chiral γ -lactam 5, easily obtained according to the procedure of Ringdahl,¹² starting with commercially available L-pyroglytamic acid (1; Scheme I). The ring opening of γ -lactam 5 was accomplished by Saito's procedure.¹¹ The optical purity of 4*R*-APA (6) thus obtained was shown to be better than 99% ee on the basis of NMR measurement of the corresponding MTPA¹³ derivative 8.

The distamycin moiety was prepared by a modification of Dervan's synthesis¹⁴ (Scheme II). Ethyl 4-nitro-1-methylpyrrole-2-carboxylate (10) was hydrogenated to give amine 11, which was coupled with γ -aminobutyric acid derivative 9 mediated by 1-ethyl-3-[3-(dimethylamino)propyl]carbodiimide (EDCI).¹⁵ The ester 12 obtained in 69% yield was saponified and again coupled with amine 11 by the EDCI method, affording the bispyrrole ester 14 in 84% yield. On the other hand, the starting ester 10 was saponified to give acid 16 in 79% yield. The subsequent peptide condensation with 3-(*N,N*-dimethylamino)propylamine (18) was accomplished by the acyl chloride method, and peptide 19 was obtained in 89% yield. The nitro group of 19 was reduced to afford amine component 20 in 89% yield. The bispyrrole acid 15, obtained in 78% yield by saponification of 14, and the monopyrrole amine 20 were coupled by the DCC-HOBt method to give the Boc-protected distamycin moiety 21 in 76% yield.

The components thus obtained were connected to each other as follows (Scheme III). *erythro*- β -Hydroxy-L-histidine (22)¹⁶ was treated with di-*tert*-butyl dicarbonate¹⁷ to give the corre-

Scheme II



sponding bis-Boc derivative 23^{4b} in 91% yield. Coupling with 4*R*-APA methyl ester (7) was effected by the DCC-HOBt procedure, furnishing dipeptide 24 in 67% yield. The Boc protection in 24 was replaced by a Z group by standard procedures in 31% yield, and a *tert*-butyl group was introduced by isobutene to give fragment 26 in 83% yield. Removal of the Z group of 26 was accomplished quantitatively with Pd-black and formic acid. Condensation of the 4-methoxypyridine moiety 28^{5c} with fragment 27 was effected with diphenyl phosphoroazidate (DPPA)¹⁸ to give 29 in 77% yield, and subsequent saponification afforded fragment 30 in 56% yield. On the other hand, the distamycin amine 31 was quantitatively prepared by treatment of the Boc derivative 21 with trifluoroacetic acid. Coupling of the amine component 31 with the acid fragment 30 was facilitated by DPPA to give the fully protected precursor 32. Conventional conditions for removal of the (*o*-nitrophenyl)thio (Nps) group¹⁹ (e.g., dilute HCl or HBr in the presence of skatole) were not successfully applied to 32 due to the facile decomposition of the distamycin residue. After extensive attempts, the deprotection was finally accomplished with 5% trifluoroacetic acid in dichloromethane to give PYML-

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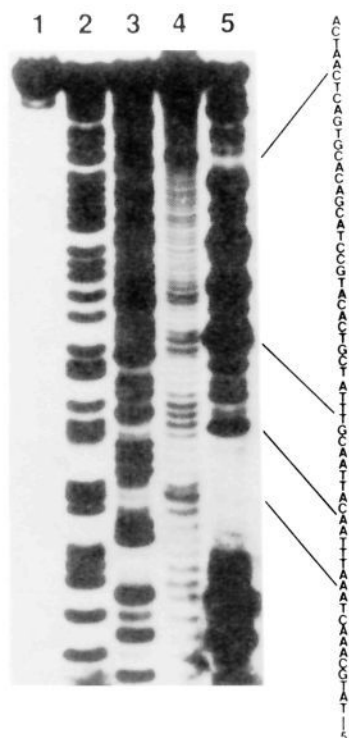


Figure 4. Nucleotide-specific cleavage of the 5'-end-labeled G4 phage DNA fragment (100 base pairs) by PYML(6)-(4R-APA)-distamycin (lane 4) and natural BLM (lane 5). Lane 1 shows the DNA blank. Lanes 2 and 3 show the Maxam-Gilbert sequencing reactions for A + G and C + T, respectively.

(6)-(4R-APA)-distamycin in 30% yield.

DNA Cleavage. Sequence-specific cleavage with PYML(6)-(4R-APA)-distamycin was examined by using a 5'-end ³²P-labeled G4 phage DNA fragment (100 base pairs) which contains AT bases in relatively high frequency. The cleavage was carried out aerobically in the presence of dithiothreitol, and the result is shown in the autoradiogram of the Maxam-Gilbert gel (Figure 4). Lanes 4 and 5 show the cleavage pattern of PYML(6)-(4R-APA)-distamycin and natural BLM, respectively. Whereas BLM mainly cleaved non-AT sites, PYML(6)-(4R-APA)-distamycin evidently cleaved A/T-rich regions preferentially.

The same G4 phage DNA fragment was labeled at the 3'-end and cleaved with PYML(6)-(4R-APA)-distamycin. AT specificity was observed in the autoradiogram of the Maxam-Gilbert gel (Figure 5). A histogram was taken from the autoradiogram, and the cleavage pattern for the 3'-end-labeled fragment was compared to that for the 5'-end-labeled strand. Figure 6 clearly shows asymmetric cleavage patterns on opposite strands, with the cleaved sites of each strand shifted to the 3' side.

Discussion

It is evident that the DNA cleavage is not an outcome of random molecular collision between BLM and DNA. The efficiency of the DNA cleavage should be regarded as a sum of both the oxygen-activating power and the magnitude of DNA affinity. DNA-cleaving activity of PYML(6)-(4R-APA)-distamycin was found to be more potent than that of PYML(6)-bleomycin, and a dramatic change in the base sequence specificity was observed. Obviously, PYML(6)-(4R-APA)-distamycin predominantly cleaved A/T-rich regions that were not cleaved by BLM (Figures 4 and 5), and the cleavage pattern is in accordance with the finding by Dervan et al. that distamycin covers 5 base pairs at A/T-rich regions.⁹ In addition, the present asymmetric cleavage pattern to the 3' side on opposite strands suggests interaction of the compound PYML(6)-(4R-APA)-distamycin with the minor groove of DNA (Figure 6).^{9b} It should also be noted that, whereas iron is placed *next* to the site of DNA binding in the DNA scission with Dervan's distamycin-EDTA-Fe(II),⁹ it appears that

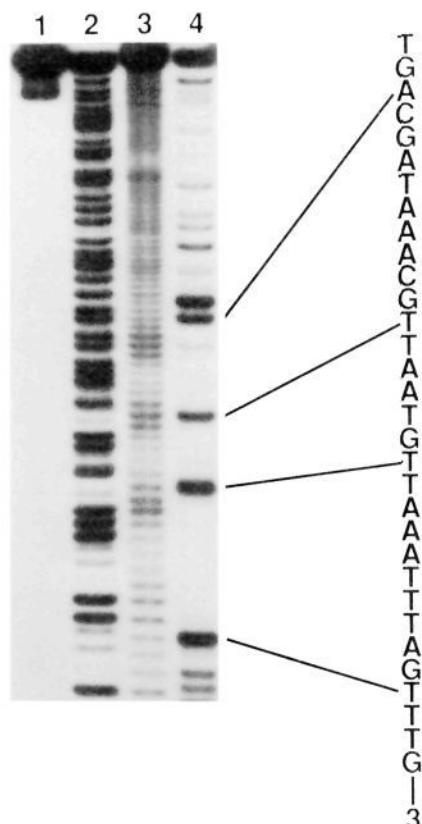


Figure 5. Nucleotide-specific cleavage of the 3'-end-labeled G4 phage DNA fragment (100 base pairs) by PYML(6)-(4R-APA)-distamycin (lane 3) and natural BLM (lane 4). Lanes 1 and 2 show the DNA blank and the Maxam-Gilbert sequencing reaction for A + G, respectively.



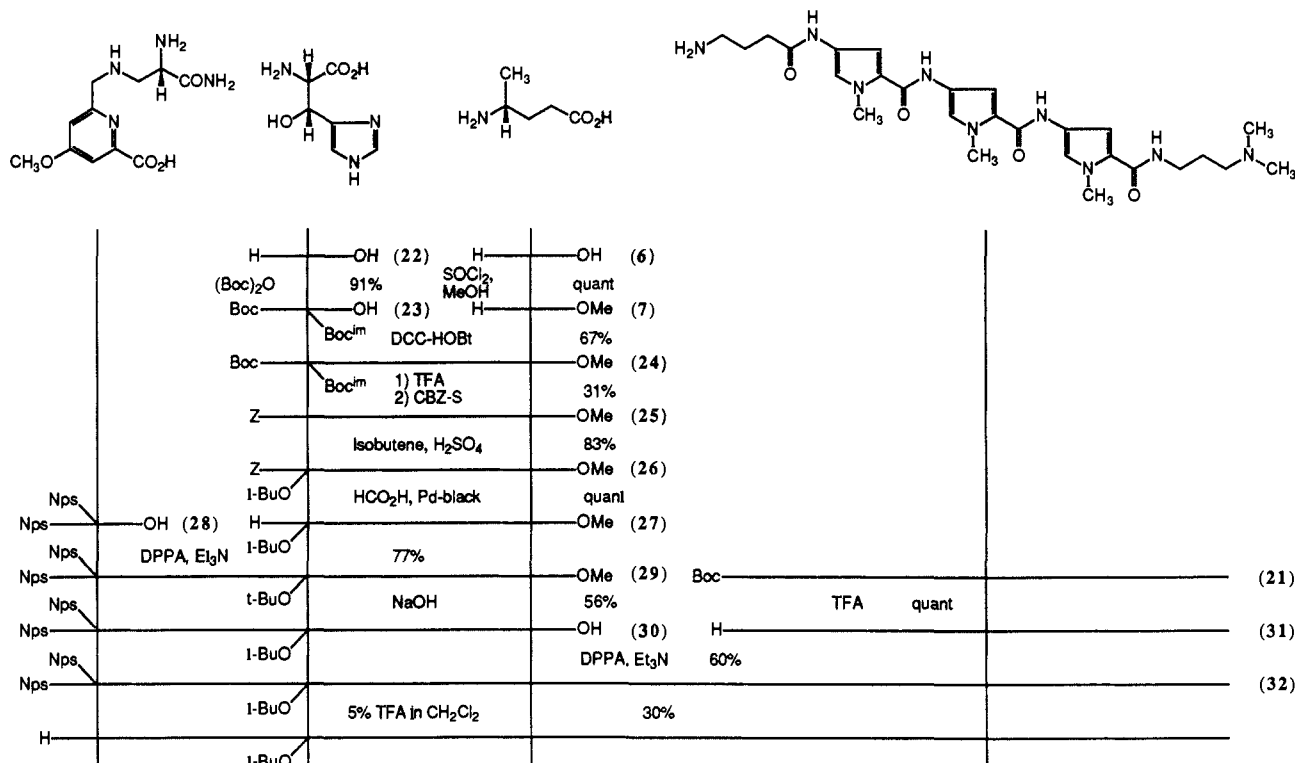
Figure 6. Histogram of cleavage sites by PYML(6)-(4R-APA)-distamycin for the G4 phage DNA fragment (100 base pairs). Relative DNA cleavage frequencies were obtained from densitometric scans of the gel autoradiograms.

PYML(6)-(4R-APA)-distamycin cleaves the site of DNA binding itself (Figure 5). This seems to be due to the conformation and the chain length of the linker moiety, which determine the placement of the metal-oxygen site relative to the location of the distamycin moiety on DNA. In fact, inspection of a molecular model of man-made BLM strongly suggests that the molecule is bending at the 4R-APA linker to orient the metal center just at the observed cleavage site. As the oxygen species activated by BLM appears to be nondiffusible and bound to iron, the sites of cleavage are indicative of the location of the iron-oxygen moiety.

Hénichart et al.²⁰ and Saito et al.²¹ independently reported simpler BLM models consisting of an ethylenediamine-pyridine-histidine chelation site, a γ -aminobutyric acid-glycine linker, and a DNA affinity site. Although nicking of supercoiled DNA by an iron complex and photochemical cleavage of DNA by a

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Scheme III



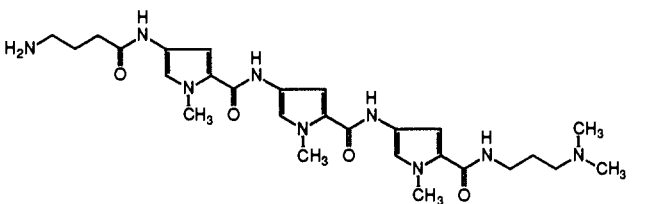
cobalt complex were demonstrated, the Maxam-Gilbert sequencing analysis showed that all bases were cleaved without any selectivity.^{20,21} Considering that the structures of these models are much simplified compared with our man-made BLMs, it appears that a highly advanced structure with complicated stereochemistry is necessary to attain specificity in DNA cleavage. We have demonstrated that the DNA affinity site with the proper linker moiety is crucial for the PYML ligand to manifest sequence specificity in DNA scission and believe that the present study provides a basis for the molecular design of novel BLM homologues with altered antitumor properties.

Experimental Section

Melting points were measured on a Yanagimoto micro melting point apparatus and are uncorrected. Optical rotations were measured on a JASCO DIP 140 instrument. ¹H NMR (400 MHz) spectra were recorded on a JEOL GX-400 spectrometer. Abbreviations are as follows: s (singlet), d (doublet), t (triplet), q (quartet), and m (multiplet). IR spectra were recorded on a JASCO DS-402G or JASCO A-102 spectrometer. Fast atom bombardment mass spectra (FABMS) were recorded on a JEOL JMS DX-300 spectrometer. Reagents and solvents were purified by standard procedures.

L-Pyrroglutamic Acid Methyl Ester (2). A solution of L-pyrroglutamic acid (1) (25.37 g, 196 mmol) in MeOH (220 mL) was cooled in a dry ice-MeOH bath. Thionyl chloride (16 mL, 221 mmol) was added over 15 min to the solution at -25 to -20 °C. The mixture was allowed to warm to room temperature and stirred for further 1.5 h. The solvent of the resulting mixture was removed in vacuo, and the residue was distilled under reduced pressure to give **2** as a colorless oil (25.07 g, 89.1%): bp 139–140 °C (2 mmHg) [lit. bp 148 °C (1 mmHg),¹² 136–138 °C (2 mmHg)²²]; IR (neat) 3230, 1735, 1690 cm⁻¹; NMR (CDCl₃) δ 2.00–2.70 (4 H, m, CH₂CH₂), 3.80 (3 H, s, CH₃O), 4.20–4.40 (1 H, m, CH), 7.13 (1 H, br s, NH).

(S)-5-(Hydroxymethyl)-2-pyrrolidone (3). NaBH₄ (6.72 g, 178 mmol) was added to an ice-cooled solution of **2** (25.04 g, 175 mmol) in EtOH (250 mL). After being stirred for 2 h at room temperature, the solution was cooled with ice, acidified with concentrated HCl, and concentrated in vacuo. The residue was suspended in CHCl₃ and filtered through celite, and the filtrate was concentrated in vacuo. The resulting pale yellow oil was purified by chromatography on silica gel (eluted with CHCl₃:MeOH = 20:1) to give **3** as colorless prisms (15.71 g, 78.0%): mp 86–87 °C (lit. mp 65–68 °C,¹² 72–73 °C,²² 66–68 °C²³); [α]_D²⁰ +34.5°



(*c* = 1.04, EtOH) [lit. [α]_D²² +29° (*c* = 1, EtOH),¹² [α]_D²⁰ +32.4° (*c* = 1.76, EtOH),²² [α]_D²⁰ +29° (*c* = 5, EtOH)²³]; IR (KBr) 3202, 1663 cm⁻¹; NMR (CDCl₃) δ 1.50–2.60 (4 H, m, CH₂CH₂), 3.10–4.00 (3 H, m, HOCH₂CH), 4.60 (1 H, br t, *J* = 5 Hz, OH), 7.50 (1 H, br s, NH).

(S)-5-(Bromomethyl)-2-pyrrolidone (4). A solution of CBr₄ (45.37 g, 137 mmol) in CH₃CN (120 mL) was added to an ice-cooled suspension of **3** (15.01 g, 130 mmol) and Ph₃P (35.88 g, 137 mmol) in CH₃CN (250 mL) under an argon atmosphere. The mixture was allowed to warm to room temperature, stirred overnight at the same temperature, and concentrated in vacuo. Water (400 mL) and hexane (400 mL) were added to the residue, and the colorless crystals deposited were removed by filtration. NaCl was added to the aqueous layer, which was extracted with CHCl₃. The extract was dried over MgSO₄ and concentrated in vacuo. The residual colorless crystals were purified by chromatography on silica gel (eluted with CHCl₃:MeOH = 20:1) to afford **4** as colorless prisms (16.89 g, 72.8%): mp 73–76 °C (lit. mp 69–72 °C,¹² 71–74 °C²³); [α]_D²⁰ -28.3° (*c* = 1.02, EtOH) [lit. [α]_D²² -32° (*c* = 1, EtOH),¹² [α]_D²² -33° (*c* = 5, EtOH)²³]; IR (KBr) 3264, 1686 cm⁻¹; NMR (CDCl₃) δ 1.70–2.70 (4 H, m, CH₂CH₂), 3.42 (2 H, d, *J* = 6 Hz, BrCH₂CH), 4.00 (1 H, distorted q, *J* = 6 Hz, CH), 7.25 (1 H, br s, NH).

(R)-5-Methyl-2-pyrrolidone (5). Tributyltin hydride (29 mL, 108 mmol) was added to a solution of **4** (16.02 g, 90.0 mmol) and AIBN (0.1414 g, 0.861 mmol) in toluene (140 mL) under an argon atmosphere. The solution was stirred at 80 °C for 5 h and concentrated in vacuo. The oily residue was purified by chromatography on silica gel (eluted with CH₂Cl₂:MeOH = 10:1). The colorless oil obtained was distilled under reduced pressure to afford **5** as a colorless oil (8.18 g, 91.7%): bp 89 °C (6 mmHg) [lit. bp 66 °C (3 mmHg)¹²]; [α]_D²² +17.2° (*c* = 1.02, EtOH) [lit. [α]_D²² +16.6° (*c* = 1, EtOH)¹²]; IR (neat) 3244, 1685 cm⁻¹; NMR (CDCl₃) δ 1.24 (3 H, d, *J* = 6 Hz, CH₃CH), 1.40–2.50 (4 H, m, CH₂CH₂), 3.81 (1 H, distorted q, *J* = 6 Hz, CH), 6.90 (1 H, br s, NH).

(R)-4-Aminopentanoic Acid (4R-APA) (6). A solution of **5** (7.88 g, 79.5 mmol) in 6 N HCl (120 mL) was heated at reflux for 8 h and concentrated in vacuo. The residue was purified by chromatography on Amberlite IR 120B (120 mL), washed with water, and then eluted with 2 N NH₃. Ninhydrin-positive fractions were collected and concentrated in vacuo to give colorless crystals that were recrystallized from MeOH, affording **6** as colorless prisms (8.44 g, 90.6%): mp 212 °C (lit. mp 209 °C¹¹); [α]_D²² +13.6° (*c* = 1.05, H₂O) [lit. [α]_D²¹ +11.8° (*c* = 1, H₂O)¹¹]; IR (KBr) 3043, 1577 cm⁻¹; NMR (D₂O) δ 1.30 (3 H, d, *J* = 7 Hz, CH₃CH), 1.60–2.50 (4 H, m, CH₂CH₂), 3.40 (1 H, sextet, *J* = 7 Hz, CH).

Methyl (R)-4-Aminopentanoate Hydrochloride (7). Thionyl chloride (0.04 mL, 0.55 mmol) was added dropwise to MeOH (1 mL) cooled with

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ice-NaCl. To the resulting solution was added **6** (20.0 mg, 0.171 mmol). The mixture was allowed to warm to room temperature, stirred for 3 days, and concentrated in vacuo to give **7** as a colorless oil (30.2 mg, quantitative): NMR (CDCl₃) δ 1.47 (3 H, distorted d, $J = 4$ Hz, CH₃CH), 1.70–2.80 (4 H, m, CH₂CH₂), 3.30–3.80 (1 H, m, CH), 3.71 (3 H, s, CH₃O), 8.45 (3 H, br s, NH₃⁺).

Methyl (*R*)-4-[*N*-[(*S*)-2-Methoxy-2-(trifluoromethyl)-2-phenylacetyl]amino]pentanoate (**8**). (*R*)-2-Methoxy-2-(trifluoromethyl)-phenylacetyl chloride¹³ (0.05 mL, 0.27 mmol) was added to a solution of **7** (30.2 mg, 0.171 mmol) in pyridine (0.5 mL) under an argon atmosphere. After the solution was stirred overnight, 3-(*N,N*-dimethylamino)propylamine (0.04 mL) and ether were successively added. The resulting mixture was washed with cold 2 N HCl, water, and saturated NaHCO₃, dried over MgSO₄, and concentrated in vacuo to afford **8** as colorless crystals (81.4 mg, quantitative): NMR (CDCl₃) δ 1.22 (3 H, d, $J = 6.6$ Hz, CH₃CH), 1.70–1.90 (2 H, m, CH₂), 2.20–2.40 (2 H, m, CH₂), 3.43 (3 H, distorted s, OCH₃), 3.65 (3 H, s, CO₂CH₃), 4.00–4.15 (1 H, m, CH), 6.78 (1 H, br d, $J = 6.6$ Hz, NH), 7.35–7.65 (5 H, m, arom).

Determination of Optical Purity of 4-Aminopentanoic Acid (**6**). Methyl (*R*)-4-[*N*-[(*S*)-2-methoxy-2-(trifluoromethyl)-2-phenylacetyl]amino]pentanoate was prepared from racemic 4-aminopentanoic acid according to the same procedure as described above: NMR (CDCl₃) δ 1.18 (3/2 H, d, $J = 6.6$ Hz, *S*-CH₃CH), 1.22 (3/2 H, d, $J = 6.6$ Hz, *R*-CH₃CH), 1.70–1.95 (2 H, m, CH₂), 2.20–2.45 (2 H, m, CH₂), 3.40 (3/2 H, distorted s, *S*-OCH₃), 3.43 (3/2 H, distorted s, *R*-OCH₃), 3.65 (3/2 H, s, *R*-CO₂CH₃), 3.70 (3/2 H, s, *S*-CO₂CH₃), 4.00–4.15 (1 H, m, CH), 6.70 (1/2 H, br d, $J = 6.6$ Hz, *S*-NH), 6.77 (1/2 H, br d, $J = 6.6$ Hz, *R*-NH), 7.35–7.65 (5 H, m, arom). The optical purity of (*R*)-4-aminopentanoic acid (**6**) prepared from L-pyrroglutamic acid (**1**) was determined to be >99% ee on the basis of the integration ratio of the 5-methyl signals.

Ethyl 4-[[4-[*N*-(*tert*-Butoxycarbonyl)amino]butyryl]amino]-1-methyl-2-pyrrolicarboxylate (**12**). A solution of ethyl 1-methyl-4-nitro-2-pyrrolicarboxylate (**10**)²⁴ (300 mg, 1.51 mmol) in DMF (10 mL) was hydrogenated over 10% Pd-C (30 mg) overnight at room temperature and atmospheric pressure. The catalyst was removed by filtration to afford a solution of crude amine **11** in DMF. 4-[*N*-(*tert*-Butoxycarbonyl)amino]butyric acid (**9**)²⁵ (305 mg, 1.50 mmol) and EDCI¹⁵ (345 mg, 1.80 mmol) were added to the solution. After being stirred for 4 h at room temperature, the solution was concentrated in vacuo. The residue was dissolved in CH₂Cl₂, washed with aqueous NaHCO₃, 5% aqueous citric acid, and saturated NaCl, dried over Na₂SO₄, and concentrated in vacuo. The residue was purified by chromatography on silica gel (eluted with CH₂Cl₂:MeOH = 20:1) to give **12** as a colorless amorphous material (365 mg, 69%): IR (CHCl₃) 3452, 1697 cm⁻¹; NMR (CDCl₃) δ 1.33 (3 H, t, $J = 7$ Hz, CH₃CH₂), 1.43 (9 H, s, CH₃C × 3), 1.70–2.10 (2 H, m, CH₂CH₂CH₂), 2.38 (2 H, t, $J = 7$ Hz, CH₂CH₂CO), 3.21 (2 H, q, $J = 7$ Hz, NHCH₂CH₂), 3.87 (3 H, s, NCH₃), 4.27 (2 H, q, $J = 7$ Hz, CH₃CH₂O), 5.35 (1 H, t, $J = 7$ Hz, NH), 6.84 (1 H, d, $J = 2$ Hz, arom), 7.38 (1 H, d, $J = 2$ Hz, arom), 9.15 (1 H, br s, NH); FABMS m/z 354 (MH⁺), 353 (M⁺).

4-[[4-[*N*-(*tert*-Butoxycarbonyl)amino]butyryl]amino]-1-methyl-2-pyrrolicarboxylic Acid (**13**). A solution of **12** (360 mg, 1.02 mmol) in MeOH (7 mL) and 1 N NaOH (2 mL) was stirred at 50 °C for 10 h. Water was added to the reaction mixture and the MeOH was removed by evaporation in vacuo. The resulting aqueous solution was cooled with ice and acidified to pH 3 with 4 N HCl. The precipitate deposited was collected and dried over P₂O₅ to give **13** as a colorless amorphous material (211 mg, 64%), which was used for the next step without further purification.

Ethyl 4-[[[4-[[4-[*N*-(*tert*-Butoxycarbonyl)amino]butyryl]amino]-1-methyl-2-pyrrolicarboxylate]amino]-1-methyl-2-pyrrolicarboxylate (**14**). A solution of **10**²⁴ (100 mg, 0.505 mmol) in DMF (3 mL) was hydrogenated over 10% Pd-C (20 mg) at room temperature and atmospheric pressure for 7 h. The catalyst was removed by filtration to afford a solution of crude amine **11** in DMF. Carboxylic acid **13** (130 mg, 0.400 mmol) and EDCI (96 mg, 0.50 mmol) were added to the solution. After being stirred overnight at room temperature, the solution was concentrated in vacuo. The residue was dissolved in CH₂Cl₂, washed with aqueous NaHCO₃, 5% aqueous citric acid, and saturated NaCl, dried over Na₂SO₄, and concentrated in vacuo. The residue was purified by chromatography on silica gel (eluted with CH₂Cl₂:MeOH = 12:1) to give **14** as a colorless amorphous material (160 mg, 84%): IR (KBr) 3422, 1685, 1654 cm⁻¹; NMR (CDCl₃) δ 1.34 (3 H, t, $J = 7$ Hz, CH₃CH₂),

1.46 (9 H, s, CH₃C × 3), 1.60–2.10 (2 H, m, CH₂CH₂CH₂), 2.36 (2 H, t, $J = 7$ Hz, CH₂CH₂CO), 3.21 (2 H, q, $J = 7$ Hz, NHCH₂CH₂), 3.90 (6 H, s, NCH₃ × 2), 4.20 (2 H, q, $J = 7$ Hz, CH₃CH₂O), 5.12 (1 H, t, $J = 7$ Hz, NH), 6.76 (1 H, d, $J = 2$ Hz, arom), 6.92 (1 H, d, $J = 2$ Hz, arom), 7.17 (1 H, d, $J = 2$ Hz, arom), 7.48 (1 H, d, $J = 2$ Hz, arom), 8.38 (1 H, br s, NH), 8.83 (1 H, br s, NH); FABMS m/z 476 (MH⁺), 475 (M⁺).

4-[[[4-[[4-[*N*-(*tert*-Butoxycarbonyl)amino]butyryl]amino]-1-methyl-2-pyrrolicarboxylate]amino]-1-methyl-2-pyrrolicarboxylic Acid (**15**). A solution of **14** (160 mg, 0.336 mmol) in EtOH (3 mL) and 1 N NaOH (0.7 mL) was stirred at 55 °C for 8 h. Water was added to the reaction mixture and the EtOH was removed by evaporation in vacuo. The resulting aqueous solution was cooled with ice and acidified to pH 3 with 4 N HCl. The precipitate deposited was collected and dried over P₂O₅ to give **15** as a colorless amorphous material (117 mg, 78%): IR (KBr) 3422, 1685, 1654 cm⁻¹; NMR (CDCl₃) δ 1.43 (9 H, s, CH₃C × 3), 1.65–2.00 (2 H, m, CH₂CH₂CH₂), 2.10–2.60 (2 H, m, CH₂CH₂CO), 3.00–3.40 (2 H, m, NHCH₂CH₂), 3.83 (3 H, s, NCH₃), 3.87 (3 H, s, NCH₃), 5.14 (1 H, br, NH), 6.88 (1 H, distorted s, arom), 6.98 (1 H, distorted s, arom), 7.11 (1 H, distorted s, arom), 7.45 (1 H, distorted s, arom), 8.45 (1 H, br s, NH), 8.81 (1 H, br s, NH); FABMS m/z 447 (M⁺).

1-Methyl-4-nitro-2-pyrrolicarboxylic Acid (**16**). NaOH (1 N, 20 mL) was added to an ice-cooled solution of **10**²⁴ (2.00 g, 10.1 mmol) in dioxane (30 mL) and EtOH (50 mL) over 10 min. After being stirred for 2 h at room temperature, the solution was concentrated in vacuo. The residue was partitioned between water and ether. The aqueous layer was cooled with ice, acidified to pH 5 with 2 N HCl, and extracted with CH₂Cl₂. The extract was dried over MgSO₄ and concentrated in vacuo to give **16** as colorless crystals (1.36 g, 79%): mp 196–199 °C (lit. 195–197 °C²⁴); IR (KBr) 3000, 1705, 1515, 1315 cm⁻¹; NMR (DMSO-*d*₆) δ 3.93 (3 H, s, NCH₃), 7.31 (1 H, d, $J = 2$ Hz, arom), 8.26 (1 H, d, $J = 2$ Hz, arom).

1-Methyl-4-nitro-2-pyrrolicarboxylic Chloride (**17**). Thionyl chloride (3.0 mL, 41 mmol) was added to **16** (1.0031 g, 5.90 mmol). The solution was heated at reflux for 3 h and concentrated in vacuo to give **17** as pale brown crystals (1.1115 g, quantitative), which was used for the next step without further purification.

N-[3-(*N,N*-Dimethylamino)propyl]-1-methyl-4-nitro-2-pyrrolicarboxamide (**19**). A solution of **17** (1.1115 g, 5.89 mmol) in CH₂Cl₂ (10 mL) was added to an ice-cooled solution of 3-(*N,N*-dimethylamino)propylamine (**18**; 0.9 mL, 7.3 mmol) and Et₃N (1.25 mL, 8.96 mmol) in CH₂Cl₂ (10 mL) over 5 min. After being stirred for 2 days at room temperature, the solution was diluted with CH₂Cl₂, washed with saturated NaHCO₃ and saturated NaCl, dried over MgSO₄, and concentrated in vacuo to give **19** as colorless prisms (1.3357 g, 89.1%): mp 129–131 °C; IR (KBr) 3300, 3140, 1655, 1500, 1305 cm⁻¹; NMR (CDCl₃) δ 1.73 (2 H, quintet, $J = 6$ Hz, CH₂CH₂CH₂), 2.33 (6 H, s, NCH₃ × 2), 2.52 (2 H, t, $J = 6$ Hz, CH₂CH₂N), 3.50 (2 H, q, $J = 6$ Hz, CONHCH₂CH₂), 4.02 (3 H, s, NCH₃), 6.98 (1 H, d, $J = 2$ Hz, arom), 7.58 (1 H, d, $J = 2$ Hz, arom), 8.63 (1 H, br, NH); FABMS m/z 255 (MH⁺).

4-[[[4-[[[4-[*N*-(*tert*-Butoxycarbonyl)amino]butyryl]amino]-1-methyl-2-pyrrolicarboxylate]amino]-1-methyl-2-pyrrolicarboxylate]amino]-1-methyl-*N*-[3-(*N,N*-dimethylamino)propyl]-2-pyrrolicarboxamide (**21**). A solution of **19** (127.1 mg, 0.500 mmol) in DMF (1.5 mL) was hydrogenated over 10% Pd-C (40 mg) at room temperature and atmospheric pressure for 2 days. The catalyst was removed by filtration, and the filtered catalyst was washed with DMF (1 mL). The DMF solutions were combined to afford a solution containing amine **20**, which was added to **15** (200.9 mg, 0.449 mmol) and HOBT·H₂O (75.8 mg, 0.495 mmol) under an argon atmosphere. The resulting solution was cooled with ice and DCC (106.7 mg, 0.517 mmol) was added. The mixture was allowed to warm to room temperature, stirred overnight, and concentrated in vacuo. The residue was purified by chromatography on silica gel (eluted with CH₂Cl₂:MeOH = 5:1) and subsequent preparative TLC (developed four times with CHCl₃:MeOH:Et₃N = 50:5:1) to give **21** as a colorless amorphous material (223.6 mg, 76%): IR (KBr) 3309, 1638 cm⁻¹; NMR (CDCl₃) δ 1.45 (9 H, s, CH₃C × 3), 1.60–2.10 (4 H, m, CH₂ × 2), 2.20–2.90 (4 H, m, CH₂ × 2), 2.30 (6 H, s, NCH₃ × 2), 3.00–3.60 (4 H, m, CH₂ × 2), 3.90 (9 H, s, NCH₃ × 3), 5.12 (1 H, br t, $J = 7$ Hz, NH), 6.68 (2 H, distorted s, arom), 6.80 (1 H, distorted s, arom), 7.11 (1 H, distorted s, arom), 7.25 (1 H, distorted s, arom), 7.30 (1 H, distorted s, arom), 7.73 (1 H, br t, $J = 7$ Hz, NH), 8.16 (1 H, br s, NH), 8.45 (1 H, br s, NH), 8.93 (1 H, br s, NH); FABMS m/z 654 (MH⁺).

N^ε,*N*^{im}-Bis(*tert*-butoxycarbonyl)-erythro- β -hydroxy-L-histidine (**23**).⁵ To an ice-cooled solution of erythro- β -hydroxy-L-histidine hydrate (197.2 mg, 1.04 mmol) in water (4 mL) were successively added Na₂SO₄ (305.9 mg, 2.89 mmol) and a solution of (Boc)₂O (557.8 mg, 2.56 mmol) in dioxane (3.2 mmol). The solution was allowed to warm to room tem-

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(6)-(4*R*-APA)-distamycin as a colorless amorphous material (9.5 mg, 30%): $[\alpha]_{D}^{22.5} + 7.9^{\circ}$ ($c = 0.365$, MeOH); IR (KBr) 3446, 1654, 1647 cm^{-1} ; NMR (CD_3OD) δ 1.16 (9 H, s, $\text{CH}_3\text{C} \times 3$), 1.24 (3 H, d, $J = 6.6$ Hz, CH_3CH), 1.65-1.90 (4 H, m), 1.90-2.10 (2 H, m), 2.15-2.40 (4 H, m), 2.92 (6 H, s, $\text{NCH}_3 \times 2$), 3.00-3.30 (4 H, m), 3.35-3.45 (2 H, m), 3.80-4.20 (6 H, m), 3.87, 3.886, 3.894, 3.91 (each 3 H, s, ArOCH_3 and $\text{NCH}_3 \times 3$), 4.60-4.70 (1 H, m), 5.26 (1 H, d, $J = 9.2$ Hz), 6.79 (1 H, d, $J = 1.8$ Hz), 6.90 (1 H, d, $J = 1.8$ Hz), 6.95 (1 H, d, $J = 1.8$ Hz), 7.06 (1 H, d, $J = 2.2$ Hz), 7.13 (1 H, d, $J = 1.8$ Hz), 7.17 (2 H, distorted s), 7.35-7.45 (1 H, m), 7.38 (1 H, d, $J = 2.6$ Hz), 7.44 (1 H, distorted s); FABMS m/z 1112 (MH^+).

DNA-Cleavage Experiment. Nucleotide sequence cleavage was investigated on the 5'- and 3'-end-labeled strands of a 100-base-pair DNA restriction fragment (*AluI-HaeIII*) from the phage R199/G4ori. The reaction mixtures contained 10 mM Tris-HCl buffer (pH 7.4), the 5'- or 3'-end ^{32}P -labeled G4 phage DNA fragment, 1 μg of carrier calf thymus DNA, 1 mM dithiothreitol, and 1 μM PYML(6)-(4*R*-APA)-distamycin (or natural BLM)-iron complex. After the reaction solutions were incubated at 37 $^{\circ}\text{C}$ for 10 min, the DNA samples were subjected to electrophoresis on a 10% polyacrylamide/7 M urea slab gel. The autoradiogram was scanned with a microdensitometer.

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Registry No. 1, 98-79-3; 2, 4931-66-2; 3, 17342-08-4; 4, 72479-05-1; 5, 21395-93-7; 6, 5937-83-7; (\pm)-6, 627-61-2; 7, 123993-04-4; (*R,S*)-8, 123993-05-5; (*S,S*)-8, 123993-18-0; 9, 57294-38-9; 10, 2853-29-4; 11, 40889-84-7; 12, 123993-06-6; 13, 123993-07-7; 14, 123993-08-8; 15, 123993-09-9; 16, 13138-78-8; 17, 28494-51-1; 18, 109-55-7; 19, 65361-30-0; 20, 78486-14-3; 21, 123993-10-2; 22, 41215-80-9; 23, 82692-03-3; 24, 123993-11-3; 25, 123993-12-4; 26, 123993-13-5; 27-HCl, 124020-63-9; 28, 108998-85-2; 29, 124020-64-0; 30, 123993-14-6; 31-TFA, 123993-15-7; 32, 123993-16-8; (*R*)-MTPA-Cl, 39637-99-5; PYML(6)-(4*R*-APA)-distamycin, 123993-17-9.

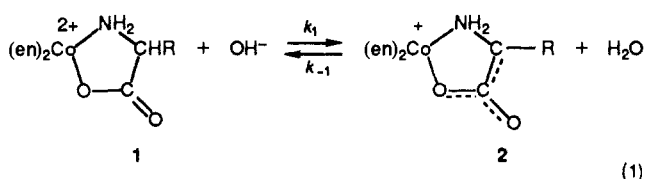
Proton Exchange and Epimerization of Co(III) Chelated Amino Acids via Carbanion Intermediates

David A. Buckingham,* Ian Stewart, and Paul A. Sutton

Contribution from the Department of Chemistry, University of Otago, Dunedin, New Zealand.
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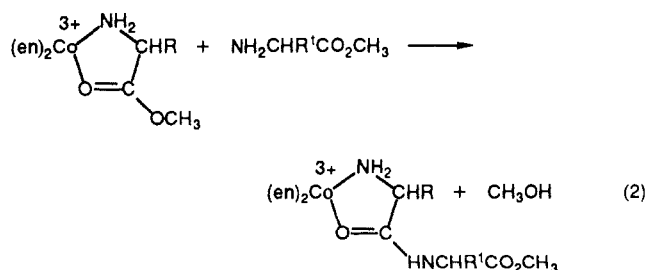
Abstract: An earlier investigation into proton exchange and epimerization of amino acids chelated to Co(III)^1 has been improved and extended to include the amino acids (AA) Phe, Val, Ala, Gly, Glu, and Asp in the complex cations $\Delta, \Delta\text{-}[\text{Co(en)}_2(\text{S-AA})]^{2+,+}$. Equilibrium concentrations of the diastereomers measured in H_2O ($K_C(\Delta\text{-S}/\Delta\text{-S}) = 1.15$ (Phe), 2.0 (Val), 1.0 (Ala, Gly), 0.85 (Glu), 0.67 (Asp)) vary little with ionic strength and are the same in D_2O . Rate constants for OD^- -catalyzed proton exchange at the 2-CH centers differ for the $\Delta\text{-S}$ and $\Delta\text{-S}$ diastereomers and can be related to the rate constant for epimerization provided the concept of a common carbanion intermediate is used. There is no correlation between the rate data and the overall charge on the complex. Selectivity differences are demonstrated in the reprotonation process ($\Delta\text{-R}/\Delta\text{-S} = 1.6$ (Phe), 1.6 (Val), 0.9 (Ala), 0.8 (Gly), 0.75 (Glu), 0.5 (Asp)), and these are shown to be thermodynamically driven. This corrects previous investigations on the AA = Asp and Glu complexes. ^3H rate studies show a kinetic isotope difference of ~ 8 for reprotonation in favor of ^1H , but no selectivity difference between ^3H and ^1H in forming the $\Delta\text{-R}$, $\Delta\text{-S}$ epimers.

This paper extends our earlier investigation into proton exchange and epimerization of chelated amino acid anions of type **1**¹ by examining in greater detail the properties of carbanion **2** generated by deprotonation of **1** by OH^- ions in aqueous solution (eq 1).



It is well-known that metal ions enhance the carbon acidity of chelated amino acids,² but the influence of the additional asymmetric metal center (structures such as **1** are diastereotopic) on the thermodynamic and kinetic stereochemical preferences for electrophilic addition to the resulting prochiral carbanion **2** is little investigated or understood.

This new study came about for two reasons. First, an investigation of epimerization during peptide synthesis using Co(III) -activated amino acid esters³ (eq 2) has revealed that car-



banions **3**, generated under the conditions of the coupling reaction, have decided diastereomeric preferences for reprotonation, and it was of interest to compare these preferences with those of the somewhat more conjugated amino acid carbanions **2**. Such information could lead to an appreciation of the steric properties, and possibly lifetimes, of such intermediates. Second, subsequent

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