

SYNTHESIS OF 3'-(L-2-AMINO-3-PHENYLPROPYL)AMINO-3'-DEOXY-N⁶,N⁶-DIMETHYLADENOSINE -
A NEW ANALOG OF PUROMYCIN

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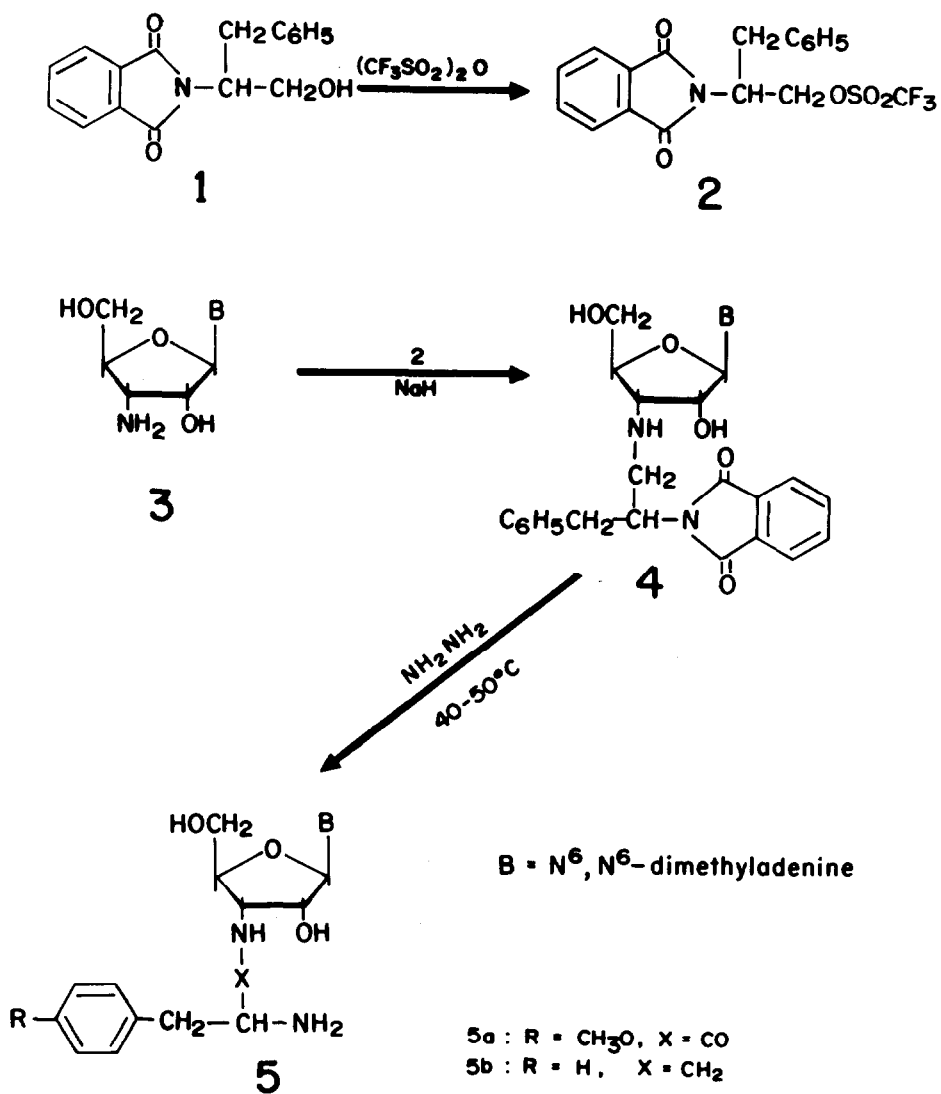
The nucleoside antibiotic puromycin (5a) is a powerful inhibitor of protein synthesis which exhibits distinct anticancer activity.¹ Analogs of puromycin (5a) wherein the carbonyl group of amino acid moiety has been reduced to a methylene group are of interest because such a modification would not increase the size of the molecule but would enhance its resistance towards chemical and enzymic hydrolysis. Such a compound would also address an important problem of the mechanism of ribosomal protein synthesis - requirement for a carbonyl group (amide or ester) in the acceptor substrate.^{2,3} Similar analogs of some peptides have found use in structure - activity studies.^{4,5}

We now wish to report the synthesis of compound (5b) using a novel selective alkylation procedure (see Scheme 1). Thus, L-3-phenyl-2-phthalimido-1-propanol (1) obtained by a method used for the corresponding racemic compound⁶ gave on reaction with trifluoromethanesulfonic anhydride in the presence of pyridine⁷ in CH₂Cl₂ trifluoromethanesulfonate 2. The interaction of 2 with puromycin aminonucleoside 3 using NaH in dimethylformamide (DMF) - dioxane mixture afforded the N-phthaloyl derivative 4. The structure of 4 was confirmed by negative ninhydrin and NaIO₄-benzidine⁸ tests. The deblocking of 4 required the treatment with hydrazine in ethanol for 5 h at 40-50°C to afford the target compound 5b characterized as a hydrochloride salt. Product 5b gave a violet coloration with ninhydrin but it did not react with NaIO₄-benzidine reagent.

Although alkylation of simple amines with esters of trifluoromethanesulfonic acid is documented⁹ the present report is the first application to sterically hindered esters and complex polyfunctional amines. The presence of trifluoromethanesulfonyloxy group as an effective leaving group in 2 is essential for the success of the procedure. Thus, the corresponding methanesulfonyl derivative of 1 failed to react with aminonucleoside 3 under a variety of conditions.

Preliminary tests in a cell-free *E. coli* ribosomal system using N-acetyl-L-phenylalanyl-tRNA as a peptide donor showed that compound 5b at concentration $1 \times 10^{-3}M$ is not capable of accepting N-acetyl-L-phenylalanyl residue from N-acetyl-L-phenylalanyl-tRNA and also it does not inhibit the puromycin reaction in the same system.¹⁰ Further biological testing is in progress.

Scheme 1



L-3-Phenyl-2-phthalimido-1-trifluoromethanesulfonyloxypropane-(2).¹¹

A solution of compound 1 (ref. 6, 0.47 g, 1.7 mmol) and trifluoromethanesulfonic anhydride (0.47 g, 1.7 mmol) in CH_2Cl_2 (10 ml) was stirred 1 h at 0°C and 1 h at room temperature. The solvent was evaporated, the residue was washed with CHCl_3 (5 ml), the solution was filtered, the filtrate was evaporated and crude 2 was crystallized from 2-propanol, 0.5 g (73%), mp. 85-86°C; NMR (CDCl_3) δ , 3.25 (m, 2, CH_2 of benzyl), 4.71 (m, 1, CH), 5.01 (m, 2, CH_2O), 7.19 (s, 5, C_6H_5), 7.73 (m, 4, phthaloyl); mass spectrum: m/e 413 (M^+).

3'-Deoxy- N^6, N^6 -dimethyl-3'-(L-3-phenyl-2-phthalimidopropyl)aminoadenosine (4).

A suspension of aminonucleoside 3 (20 mg, 68 μmol) in DMF-dioxane (1:1, 4 ml) was stirred with NaH (2 mg, 83 μmol) for 30 min. at room temperature. Reagent 2 (28 mg, 68 μmol) was then added and the stirring continued for 17 h. The insoluble portion was filtered off, the filtrate was evaporated and the residue was chromatographed on a loose layer of silica gel¹² in CH_2Cl_2 - CH_3OH (93:7). The faster moving band was eluted, the eluate was evaporated and crude 4 was crystallized from 2-propanol or ethyl acetate, 21 mg (55%), mp. 202-203°C; $\text{UV}_{\text{max}}^{\text{EtOH}}$ 267 nm (ϵ 17 300), shoulder 283 nm (ϵ 15 600); NMR ($\text{CD}_3\text{SOCD}_3 + \text{D}_2\text{O}$) δ , 3.50 (s, 6, $(\text{CH}_3)_2\text{N}$), 6.11 (d, $\text{J}_{1',2'}$, 3 Hz, 1, $\text{H}_{1'}$), 7.18 (s, 5, C_6H_5), 7.73 (m, 4, phthaloyl), 8.31 (s, 1, H_2), 8.46 (s, 1, H_8); mass spectrum: m/e 557 (M^+). $[\alpha]_{\text{D}}^{24} - 6.2^\circ$ (c 0.15, ethanol).

3'-(L-2-Amino-3-phenylpropyl)amino-3'-deoxy- N^6, N^6 -dimethyladenosine (5b).

A mixture of compound 4 (20 mg, 36 μmol), hydrazine hydrate (1 ml) and ethanol (4 ml) was heated for 5 h at 40°-50°C. After evaporation of solvents the residue was chromatographed on a loose layer of silica gel¹² in CH_2Cl_2 - CH_3OH (4:1). The major UV absorbing band was eluted, the eluate was evaporated, product was dissolved in methanol (20 ml), the solution was saturated with HCl and then it was kept at -10°C for 3 days. The insoluble portion was filtered off, the filtrate was evaporated and the residue was crystallized from 80% ethanol, 11 mg (66%, hydrochloride); mp. 223-227°C; $\text{UV}_{\text{max}}^{\text{EtOH}}$ 275 nm (ϵ 13 600); NMR ($\text{CD}_3\text{SOCD}_3 + \text{D}_2\text{O}$) δ , 3.46 (s, 6, $(\text{CH}_3)_2\text{N}$), 5.99 (d, $\text{J}_{1',2'}$, 3 Hz, 1, $\text{H}_{1'}$), 7.25 (s, 5, C_6H_5), 8.21 (s, 1, H_2), 8.40 (s, 1, H_8). $[\alpha]_{\text{D}}^{24} - 9.8^\circ$ (c 0.12, H_2O).

This research was supported in part by Grant CA 21388 from the National Cancer Institute, Bethesda, Maryland and in part by an institutional grant from the United Foundation of Greater Detroit. The authors are indebted to Dr. H.L. Chung, D. Andrzejewski and W. Brukwinski for spectroscopic measurements.

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11. All new compounds gave satisfactory C, H, N analyses and they were homogeneous on TLC.¹² NMR spectra were measured on FT spectrometer JEOL FX 100 and mass spectra on JEOL JMS 01SG-2. Tetramethylsilane was used as an internal standard with CDCl_3 and sodium 2, 2-dimethyl-2-silapentane-5-sulfonate with CD_3SOCD_3 . Evaporations were carried out on a rotary evaporator at 30°C. Melting points were determined on a Thomas-Hoover capillary melting point apparatus and are not corrected.
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(Received in USA 8 August 1978)