## A Circular Dichroic Method for determining the Linkage to Symmetric Hexocyclitols and Hexocyclitolamines: Structure of Two Antibiotics from *Nocardia* sp.

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Summary A highly sensitive c.d. method which appears to have general applicability has been developed to determine the point of attachment of substituents on symmetric hexocyclitolamine and hexocyclitol residues; it has been applied in the structure determination of two antibiotics LL-BM 123 $\alpha$  and LL-BM 782 $\alpha_2$ .

HEXOCYCLITOLAMINE and hexocyclitol residues are present in a number of natural products such as the antibiotics (1)and (9) (thick lines). Because of the symmetry of these moieties, no chemical method has yet been described which resolves the point of attachment of other groups to available enantiotopic hydroxy-groups such as at positions 4 and 6 on the cyclitol subunits of (1) and (9). X-Ray analysis of course provides a solution but often these natural products, even in a highly purified form, are very difficult to crystallize. We describe here a practical and highly sensitive c.d. method which appears to have general applicability to solve such problems.



The antibiotic LL-BM 123 $\alpha$  isolated from a species of *Nocardia* exhibits moderate *in vivo* activity against gramnegative bacteria and is remarkably nontoxic; structural studies led to the structure (1) where it was left undecided whether the mannose unit was linked to 4" or 6" (arrows) of the myoinosamine unit (and with the  $\alpha$ -configuration at 1').<sup>1</sup> The following evidence leads to the full structure (1). The trisaccharide (2)<sup>†</sup> (25 mg)<sup>1</sup> was *N*-acetylated with

MeOH-Ac<sub>2</sub>O (75 min, room temp.) to give (3), † which was permethylated (NaH-Me<sub>2</sub>SO-MeI) to afford 9 mg of (4).† Methanolysis (MeOH-HCl, 1:1 v/v, 5 h) of (4) and column chromatography gave 4 mg of the N-acetyl-O,N-permethylated glucosaminylmannose derivative and four other products as detectable spots on t.l.c. with common spraying agents. However, since none of them showed a <sup>1</sup>H n.m.r. spectrum corresponding to the expected product (5), the remaining eluates were combined and benzoylated for spectroscopic detection; column chromatography and t.l.c. of the benzoylated eluate gave  $0.5 \text{ mg of } (\mathbf{6})$ ,  $\dagger$  which was hydrolysed to (7) by dissolving it in a trace of MeOH and refluxing in aqueous Ba(OH)<sub>2</sub> for 3 h; (7)<sup>†</sup> showed OMe <sup>1</sup>H n.m.r. signals at  $\delta$  3.59, 3.49, 3.44, and 3.42 and one NMe signal at  $\delta$  2.57. Treatment of (7) with *p*-bromobenzoyl chloride in pyridine-Et<sub>a</sub>N at 0 °C overnight gave 50 µg of the p-bromobenzoate (8)<sup>†</sup> which was purified by column chromatography and t.l.c.

Chemical ionization mass spectrometry with methane confirmed (8) to be the O,N-dibenzoyl compound. The amount of this compound, in a cell, was estimated from the standard u.v.  $\epsilon$  value of 38,200 at 244.5 nm for di-pbromobenzoates<sup>2</sup> without weighing of the sample, and from this it was possible to estimate<sup>‡</sup> the  $\Delta \epsilon$  value of the excitonsplit c.d. curve at 253/236 nm.<sup>3,4</sup>

Since (8) exhibits a positive split c.d. with an amplitude of +28,§ the OBz group is located at the 4"- and not the 6"-position. The point of attachment to the mannose unit is therefore also the 4"-position. In BM  $123\alpha$   $^{1}J_{CH}$  for C-1' is 162.2 Hz. From the published values for  $\alpha$ - and  $\beta$ -anomers, the  $^{1}J_{CH}$ <sup>5-7</sup> value shows that the anomeric configuration at 1' is  $\beta$ . This leads to structure (1) for this antibiotic.



Structure (9) was assigned to the broad-spectrum antibiotic LL-BM  $782\alpha_2$ ,<sup>8</sup> where an  $\alpha$ -configuration was tentatively assigned to C-1 and attachment of the 3-deoxy-3-guanidino-D-mannose to myoinositol was left open (4' or 6', arrows).

<sup>†</sup> Compound which was purified by column chromatography and characterized by <sup>1</sup>H n.m.r. and mass spectroscopy, etc.

<sup>‡</sup> The u.v. and c.d. properties of N-benzoates are similar to those of O-benzoates.

§ The difference in  $\Delta \epsilon$  values between the longer and shorter wavelength extrema of split Cotton effects.

The acid hydrolysis product (10)<sup>9</sup><sup>†</sup> (10 mg) was permethylated to give (11),<sup>†</sup> the <sup>1</sup>H n.m.r. spectrum of which showed six OMe signals ( $\delta$  3.68–3.40) and five NMe signals  $(\delta 3.01-2.20)$ . Permethylated (11) was submitted to methanolysis (20 h), Ba(OH), hydrolysis (18 h) and p-bromobenzoylation to yield (12).<sup>†</sup> The above-mentioned m.s.  $\rightarrow$ u.v.  $\rightarrow$  c.d. sequence gave  $\Delta\epsilon_{252} + 35\cdot 3/\Delta\epsilon_{235} - 15\cdot 9$  or A = +51. In the 1',3',4'- (or 1',3',6'-) tribenzoate system (12), the dibenzoate systems 1',3' and 1',4' (or 6') do not lead to chirality. The positively split Cotton effects (+51) unambiguously show that the chirality is between the 3',4'dibenzoate system, and that therefore the original linkage is at 4'. The  ${}^{1}J_{CH}$  163.9 Hz (C-1) value again shows that the anomeric linkage is  $\beta$ .

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¶ Location of N-Me groups unassigned.

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