## SYNTHESIS OF THE LIPOSIDOMYCIN DIAZEPANONE

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Abstract: Acrolein and two sarcosines are combined to give the diazepanone substructure of the liposidomycin nucleoside antibiotics. Conformational analysis of the substituted diazepanone ring is presented.

The liposidomycins are complex, lipid-bearing nucleoside antibiotics (e. g. 1 and 2) that selectively inhibit bacterial peptidoglycan synthesis.<sup>1,2</sup> Liposidomycin C (2) has been shown to specifically inhibit *in vitro* the phospho-N-acetylmuramyl-pentapeptide transferase from *E. coli* Y-10, which catalyzes the first step in the peptidoglycan lipid cycle.<sup>3</sup> Structures were ascribed to 1 and 2 based on degradation studies and NMR and mass spectral evidence, but several stereogenic centers on or near the diazepanone ring (5', 6', 2''', and 3''') remain unassigned.<sup>2</sup> As a step toward elucidating the relative and absolute stereochemistry of 1 and 2, we report the synthesis of the diazepanone substructure as represented by 3 and 4, and also the conversion of 3 and 4 to tetrahydrodiazepinone 13, a substructure of the products of acid and base hydrolysis of 1.



Aldol reaction of the anion of 5 and acrolein gave adduct 6 as a 4:1 anti/syn mixture of diastereomers. Because 6 is not stable to chromatography, it was acylated directly and isolated from the crude reaction mixture as the benzoate 7. Removal of the BOC and neutralization led to a 7:1 mixture of amines 8; the apparent enrichment in the anti diastereomer is the result of selective loss of the syn diastereomer by an O-to-N benzoyl migration. IIDQ-mediated coupling<sup>4</sup> with Z-sarcosine gave the dipeptides 9 and 10, easily separable at this stage, along with a small amount of the benzoyl migration product. The stereochemistry of the aldol adducts 6 was proven by in situ lithium borohydride reduction to a 4:1 mixture of diols, each of which was purified and separately converted to an acetonide with diagnostic proton coupling constants.<sup>5</sup> The predominance of the anti addol product from the lithium anion of 5 is consistent with preferential (and typical<sup>6</sup>) formation of the E ester enolate isomer. Ozonolysis of the vinyl group and reductive cyclization gave the trans diazepanone 3 from purified anti dipeptide 9, and cis diazepanone 4 from purified syn dipeptide 10. Fully decoupled <sup>1</sup>H NMR spectra of 3 and 4 suggest the solution conformations shown. In particular, the H(6)-H(7) coupling constants (J = 5.2 and 2.2, respectively) are consistent with pseudo-axial carbethoxy substituents, which avoid a repulsive interaction with the N(1) methyl group. A five-bond "W" coupling (J = 2) between H(3<sub>eo</sub>) and  $H(5_{cn})$  is observed for 3. These conformations are reproduced as minima by the MacroModel<sup>®</sup> program (Amber subroutine),<sup>7</sup> which also predicts<sup>8</sup> H(6)-H(7) coupling constants of 5.1 and 1.8, respectively.





The *trans* and *cis* diazepanones 3 and 4 were each converted to the tetrahydrodiazepinone ester 11 under mild conditions by treatment with DBU. Hydrolysis of the ester group gave the carboxylate salt 12, and protonation led to the ammonium salt 13. The ability to generate this tetrahydrodiazepinone carboxylate ring system from any of several substituted diazepanone esters should be useful for the synthesis of the liposidomycin degradation<sup>2</sup> product 14, and thus assignment of the stereochemistry at C(5') and C(6') of 1.



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