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ADDITIONS AND CORRECTIONS

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Complexation and medium effects on the conformation of cyclosporin A studied by NMR spectroscopy and molecular dynamics calculations

Horst Kessler, Matthias Gehrke, Jörg Lautz, Matthias Köck, Dieter Seebach and Adrian Thaler

Pages 170–172, Figures 1 through 4 were printed with the incorrect legends. The figures and legends should have appeared as follows:

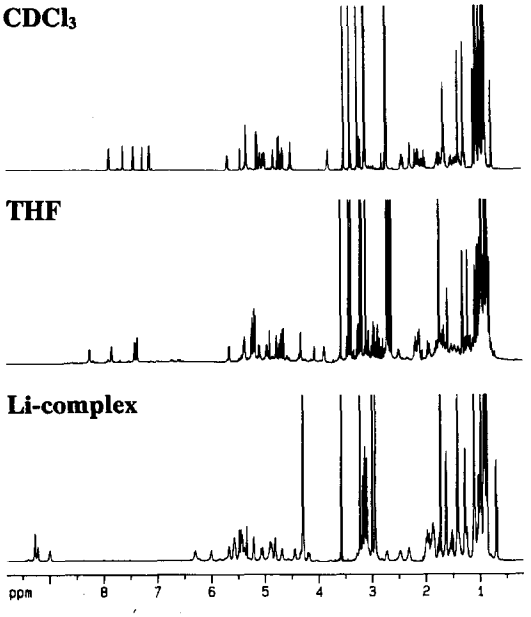

Fig. 1. 600 MHz ^{1}H NMR spectra of cyclosporin A in $CDCl_3$, THF-d_8 and in THF-d_8 with addition of 30.9 equivalent LiCl.

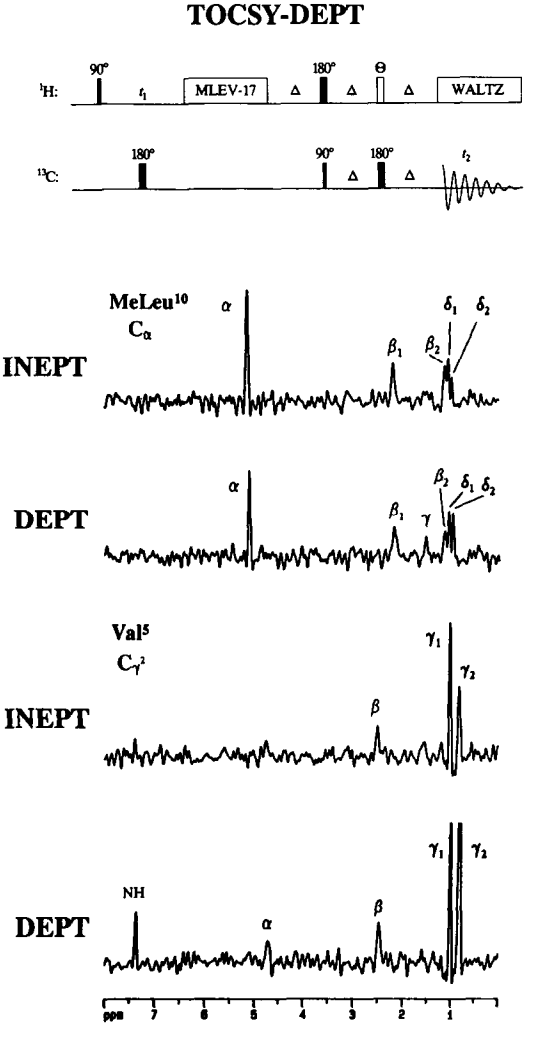

Fig. 2. Cross-sections of C_α of MeLeu^{10} and $C_{\gamma 2}$ of Val^5 of the 250 MHz TOCSY-DEPT spectrum of CSA in THF-d_8. For a comparison, the analogue TOCSY-INEPT spectrum is shown. The total spin systems of the amino acid proton spin systems of MeLeu^{10} and Val^5 can be assigned on the chemical shift of the two carbons.

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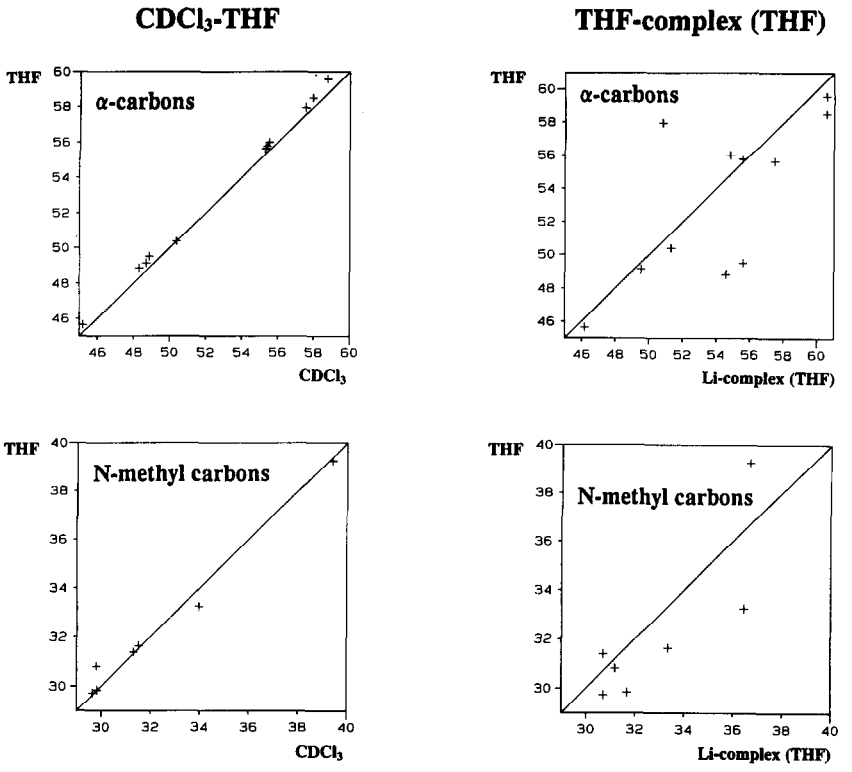

Fig. 3. Comparisons of the chemical shifts of the α-carbons and the N-methyl carbons of CSA in $CDCl_3$, THF (left) and THF without and with lithium salt addition (right).

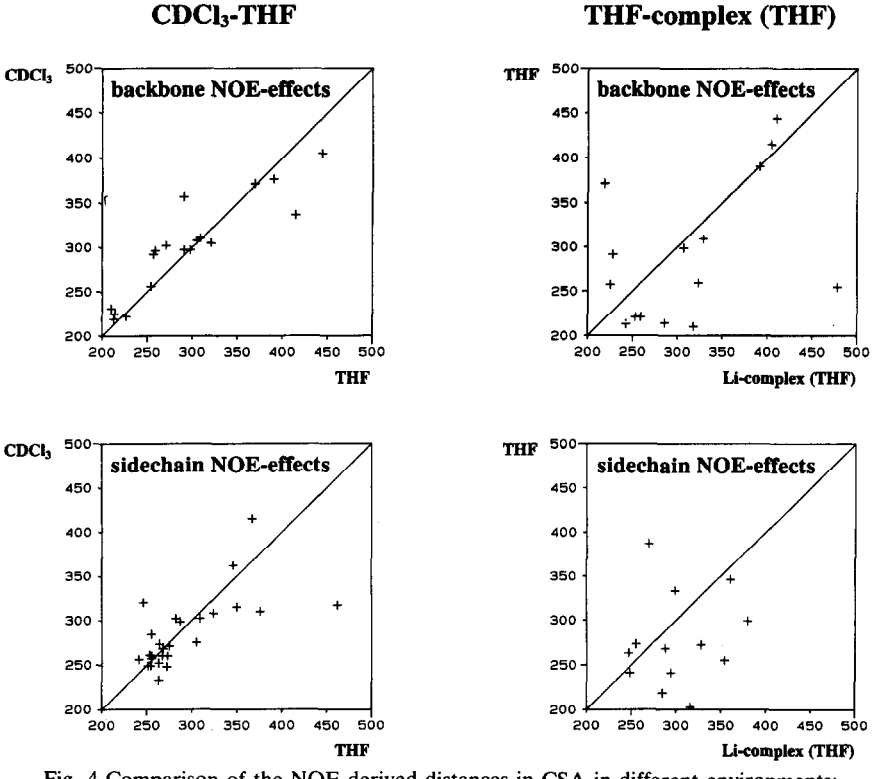

Fig. 4 Comparison of the NOE-derived distances in CSA in different environments: $CDCl_3$/THF and THF without and with addition of lithium salt.