

## Spatial structure of peptides determined by residual dipolar couplings analysis

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### Abstract

The gated decoupled  $^{13}\text{C}$  NMR spectra of a dipeptide (Glu-Trp) and a tetrapeptide (NAc-Ser-P-e-Val-Gly-OMe) were recorded in  $\text{D}_2\text{O}$  and in a lyotropic alignment medium (pentaethylene glycol monododecyl ether/*n*-hexanol). The residual dipolar couplings were extracted as the differences between the observed couplings for the magnetic nuclei dissolved in the latter and former media. Using a computational optimization, the spatial structures of the compounds were calculated starting from their respective low energy conformations obtained on a semiempirical basis. The uniformity of each conformation was confirmed by the solid-state  $^{13}\text{C}$  NMR spectra of powder samples. Differences between the starting structures and final ones, optimized when employing residual dipolar couplings, are discussed. Copyright © 2008 John Wiley & Sons, Ltd.

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### Keywords

$^{13}\text{C}$  NMR, Lyotropic medium, NMR, Oligopeptides, Pentaethylene glycol monododecyl ether (C12E5), Residual dipolar couplings, Spatial structure