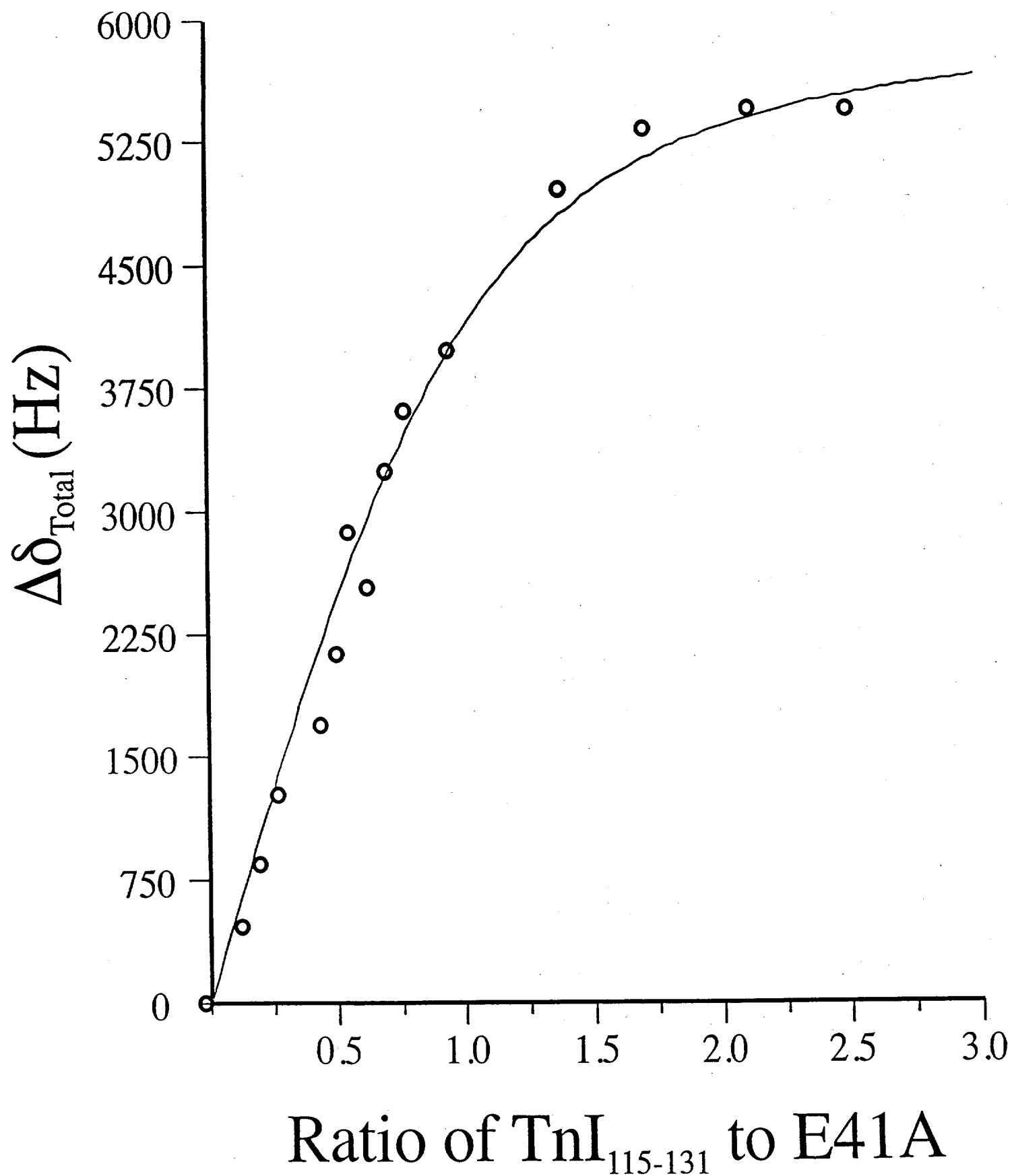


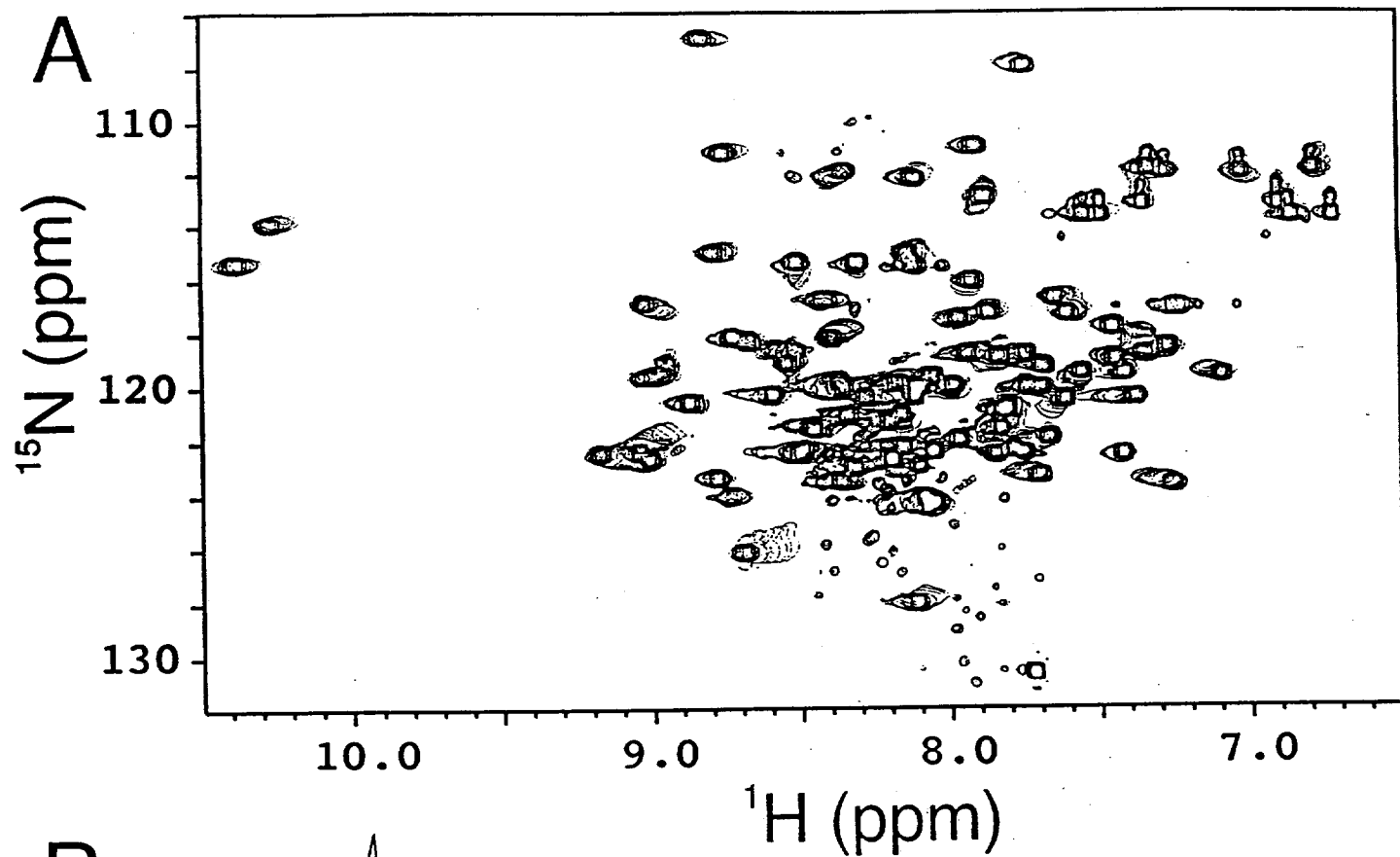


Figure 1S. Binding curve determined from the $\Delta\delta_{\text{total}}$ of 2D $\{^1\text{H}, ^{15}\text{N}\}$ -HSQC NMR spectral changes of E41A-sNTnC•2Ca²⁺ upon addition of sTnI₁₁₅₋₁₃₁. The curve is an average of 82 residues, which exclude Ser¹⁵, Asp³⁰, Asp³⁶, Val⁶⁵, and Glu⁷⁶ due to spectral overlap, Pro⁵³ due to the lack of an amide hydrogen resonance, Ala¹ and Ser² presumably due to rapid exchange with solvent, and residues involved in site I. The determined dissociation constant was 300 ± 100 μM .

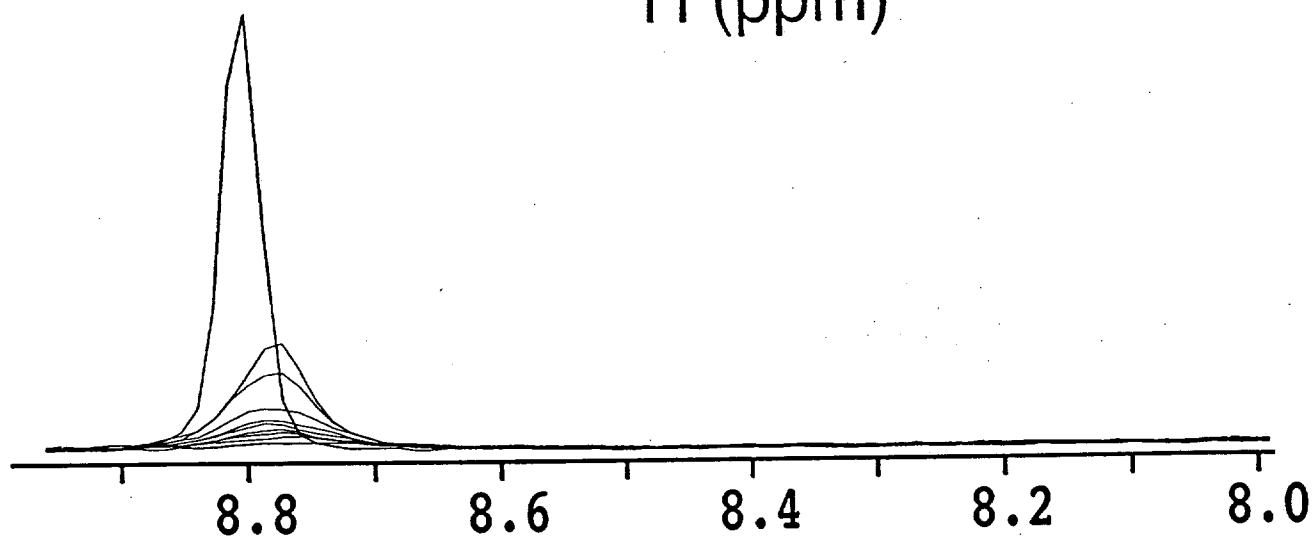
Figure 2S. (A) Contour plot of 2D $\{^1\text{H}, ^{15}\text{N}\}$ -HSQC NMR spectra of apo E41A-sNTnC showing the effect of sTnI₁₁₅₋₁₃₁ addition. The apo E41A-sNTnC spectrum before the addition of peptide is shown with multiple contours while each addition of sTnI₁₁₅₋₁₃₁ are indicated by a single contour. (B) 1D-trace of Gly⁴³ throughout the titration of sTnI₁₁₅₋₁₃₁, showing the progressive line broadening effect.

Figure 3S. The total change in chemical shift ($\Delta\delta_{\text{total}}$ in Hz) for each backbone amide pair in E41A-sNTnC (A,D), cNTnC (B,E) and sNTnC (C,F) upon addition of calcium (A-C), or TnI peptide (D-F). The thin dashed lines indicate one standard deviation above the average.

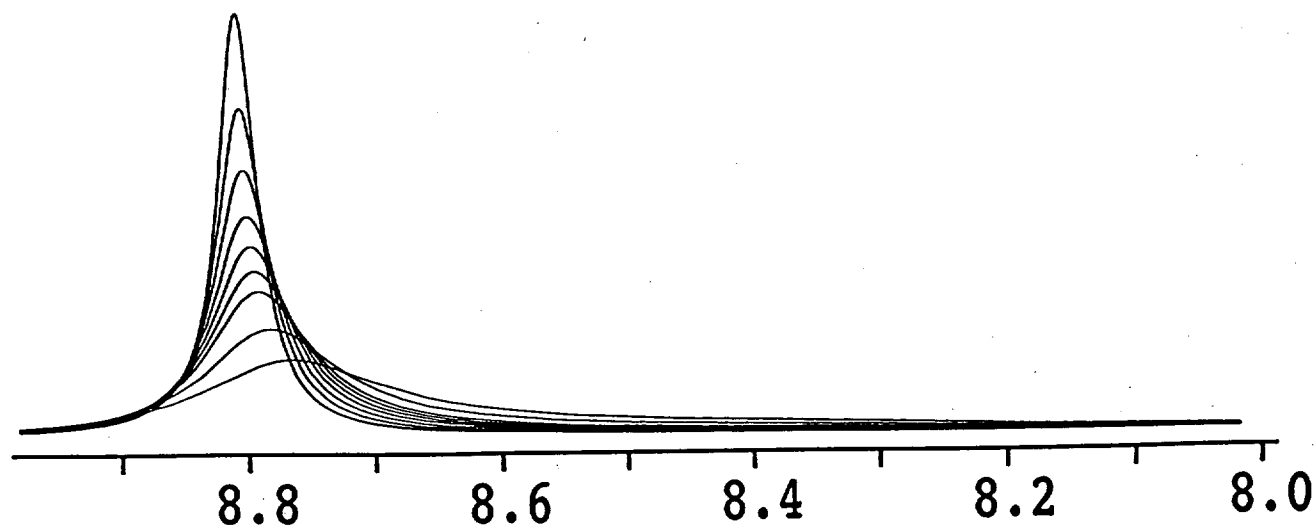


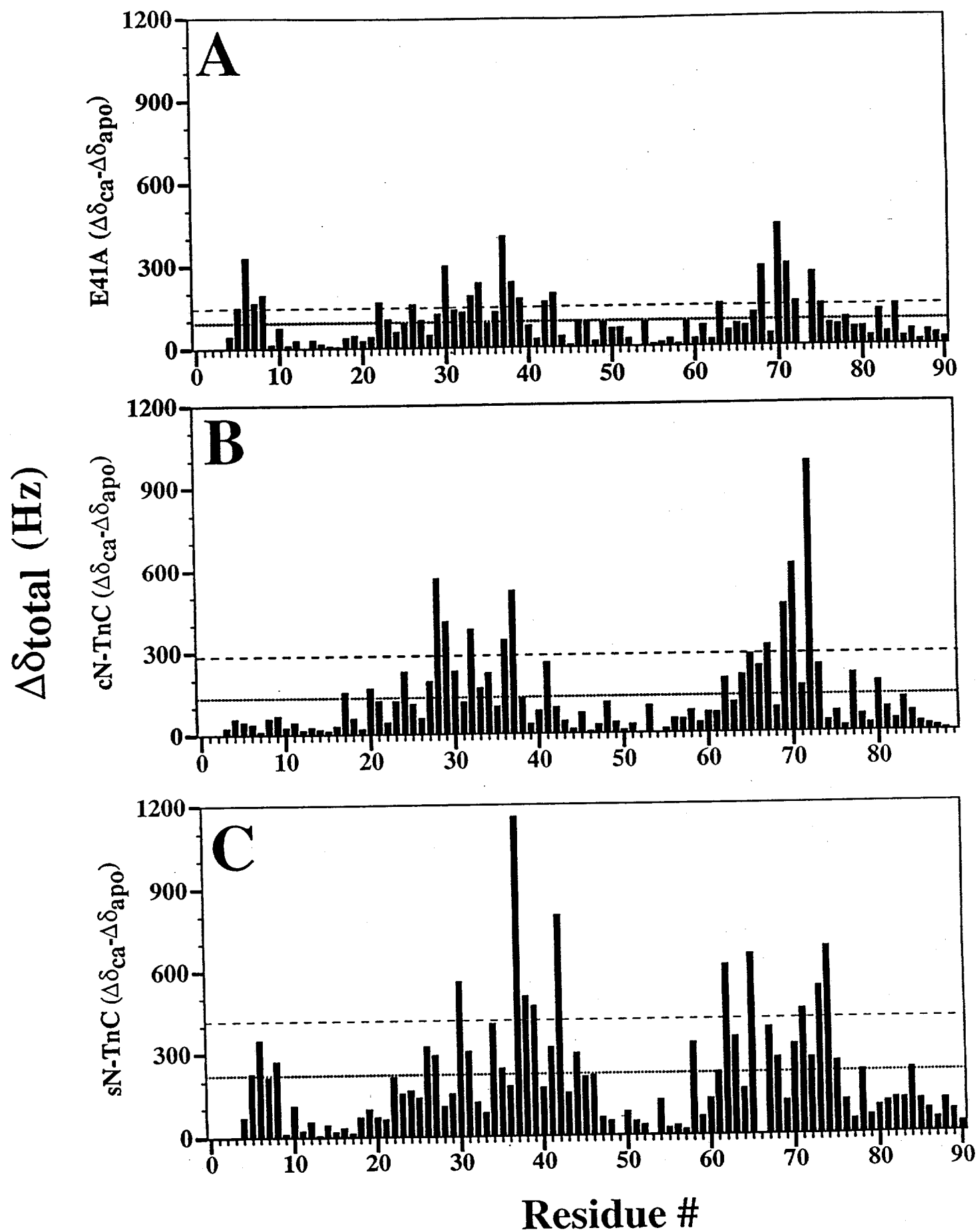


B



C





$\Delta\delta_{\text{total}}$ (Hz)

