

Correction to “UV Resonance Raman Study of TTR(105–115) Structural Evolution as a Function of Temperature”

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The legend of Figure 9 was published incorrectly. The corrected Figure 9 is as follows:

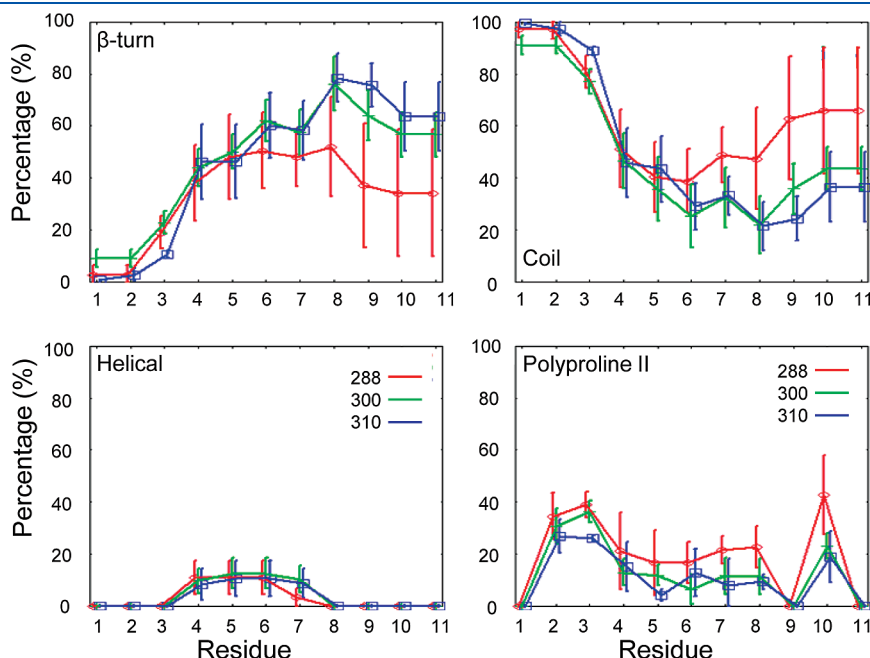


Figure 9. Secondary-structure probability of the peptide residues, computed from the simulations at 288, 300, and 310 K simulations. The probabilities for occupation of polyproline II conformation presented in the bottom right panel are also included in the random coil probabilities in the upper right panel. The standard deviation is shown in error bars and was calculated using the average probability populations of five successive windows of 1 ns duration each. Residues 1–11 correspond to segment 105–115 of the TTR sequence.

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