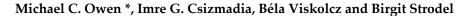
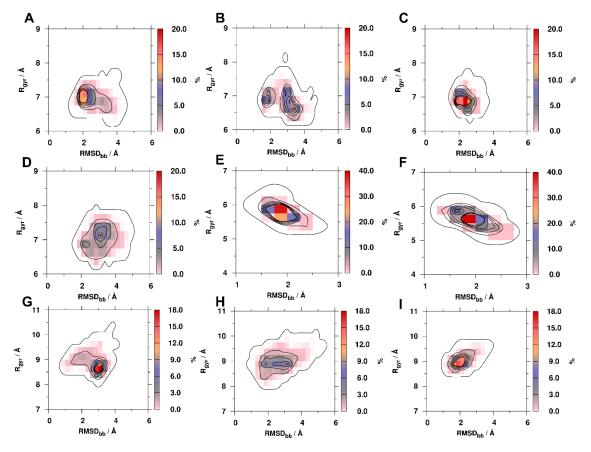
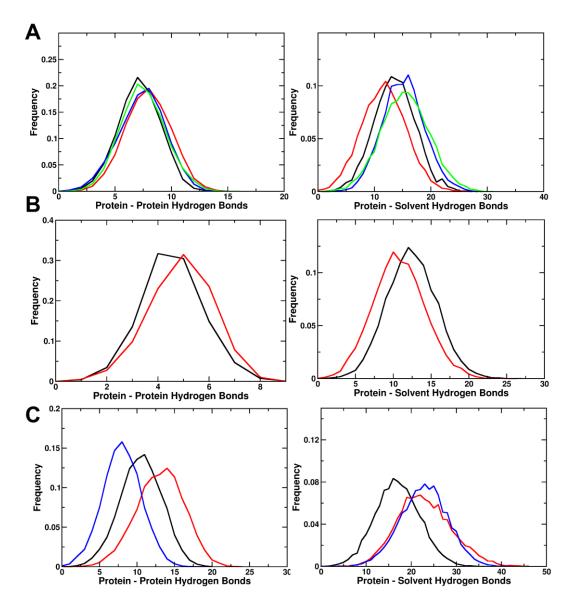
## Supplementary Materials: Protein Stability and Unfolding Following Glycine Radical Formation





**Figure S1.** The Rgyr-RMSD density plots of the Trp cage ( $\mathbf{A}$ ), Trp cage  $C_{\alpha}$ -centered Gly radical in place of Gly10 (GLR10) ( $\mathbf{B}$ ), Trp cage(GLR11) ( $\mathbf{C}$ ), Trp cage(GLR15) ( $\mathbf{D}$ ), Trp zipper ( $\mathbf{E}$ ), Trp zipper(GLR6) ( $\mathbf{F}$ ), the villin headpiece ( $\mathbf{G}$ ), villin(GLR11) ( $\mathbf{H}$ ) and villin(GLR33) ( $\mathbf{I}$ ).

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**Figure S2.** The number of protein-protein hydrogen bonds (left panel) and protein-solvent hydrogen bonds (right panel). Row **A** contains the data for the Trp cage (in black), Trp cage(GLR10) (in red), Trp cage(GLR11) (in red), Trp cage(GLR15) (in blue). Row **B** contains data for the Trp zipper (in black), Trp zipper(GLR6) (in red). Row **C** contains data for the villin headpiece (in black), villin(GLR11) (in red) and villin(GLR33) (in blue).