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Protein Folding Kinetics

Biophysical Methods

With 145 Figures and 15 Tables

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Cover illustration: Structure of the 10-kDa protein barstar, and structural characteristics of its microsecond folding transition state (Nölting et al., 1997a; Nölting, 1998a). Amino acid residues which have a significant Φ-value are highlighted with a ball-and-stick structure. The figure was generated using MOLSCRIPT (Kraulis, 1991), Raster3D (Bacon and Anderson, 1988; Merritt and Murphy, 1994) and the NMR structure of barstar (Lubienski et al., 1994).


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