

Corrections

Hierarchical Map of Protein Unfolding and Refolding at Thermal Equilibrium Revealed by Wide-Angle X-ray Scattering, by Mitsuhiro Hirai,* Masaharu Koizumi, Tomohiro Hayakawa, Hiroshi Takahashi, Satoru Abe, Harutaka Hirai, Keiko Miura, and Katsuaki Inoue, Volume 43, Number 28, July 20, 2004, pages 9036–9049.

Page 9046. Figure 10 was published without a color legend. The correct version follows.

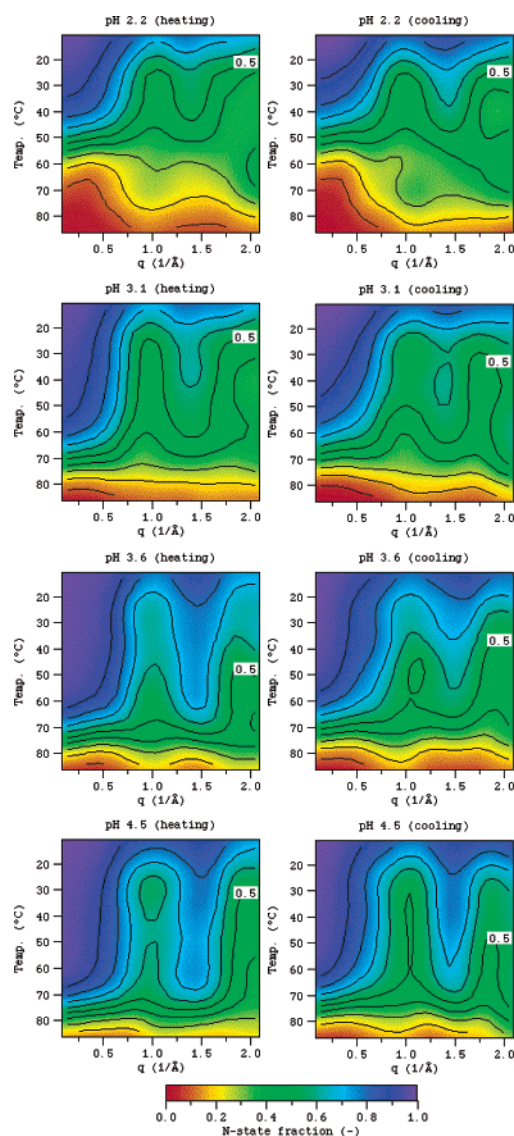


FIGURE 10: Two-dimensional hierarchical maps of unfolding–refolding transitions of the HEWL structure against temperature and q value, termed SH maps, which are obtained from the molar fractions α of native-like structures for different q ranges (q intervals of 0.05 \AA^{-1}) in the WAXS curves in Figure 4. The α values are displayed with rainbow colors and contour lines. The $\alpha = 0.5$ contour line mostly corresponds to the transition midpoint for each hierarchical structure.

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