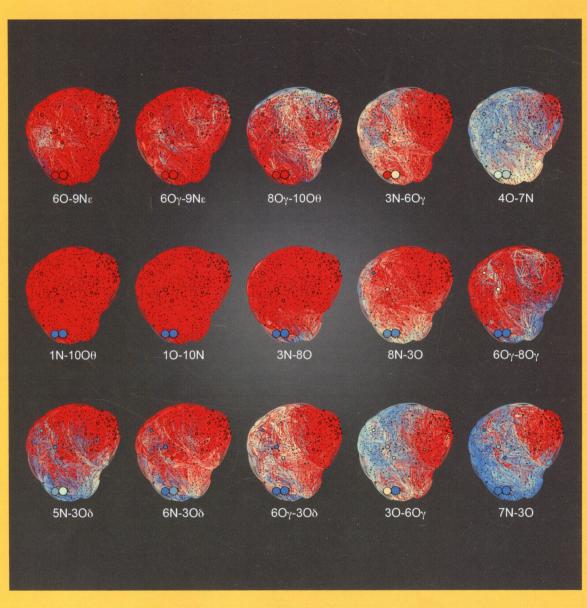
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# THE JOURNAL OF PHYSICAL CHEMISTRY

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Configuration Space Networks of Chignolin Reveal How Hydrogen Bonds Guide Folding (see page 5A)

BIOPHYSICAL CHEMISTRY, BIOMATERIALS, LIQUIDS, AND SOFT MATTER



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**ON THE COVER:** Configuration space networks of chignolin reveal how hydrogen bonds guide folding. In Dickson et al., an algorithm ("WExplore") is introduced that enhances sampling and allows for the efficient construction of configuration space networks. In CSNs, each node (circle) represents a configuration of the system, and the edges (lines) join together configurations that are connected by transitions during sampling. The native state is composed of the two large nodes at the bottom left, and the top right states are the most unfolded. The nodes on the cover are colored according to the distance between various native and nonnative hydrogen bonding pairs, with blue indicating the pairs are within hydrogen-bonding distance, and red indicating they are far away. This visualization technique allows for the easy identification of nonnative interactions that help guide folding (such as 40–7N) as well as native hydrogen bonds that are formed early (7N–3O) or late (10–10N) in the folding process. See page 3532.

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3498 dx.doi.org/10.1021/jp4097327

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3507 dx.doi.org/10.1021/jp410477x

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Marieke Füllbrandt, Elena Ermilova, Asad Asadujjaman, Ralph Hölzel, Frank F. Bier, Regine von Klitzing, and Andreas Schönhals\*

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