

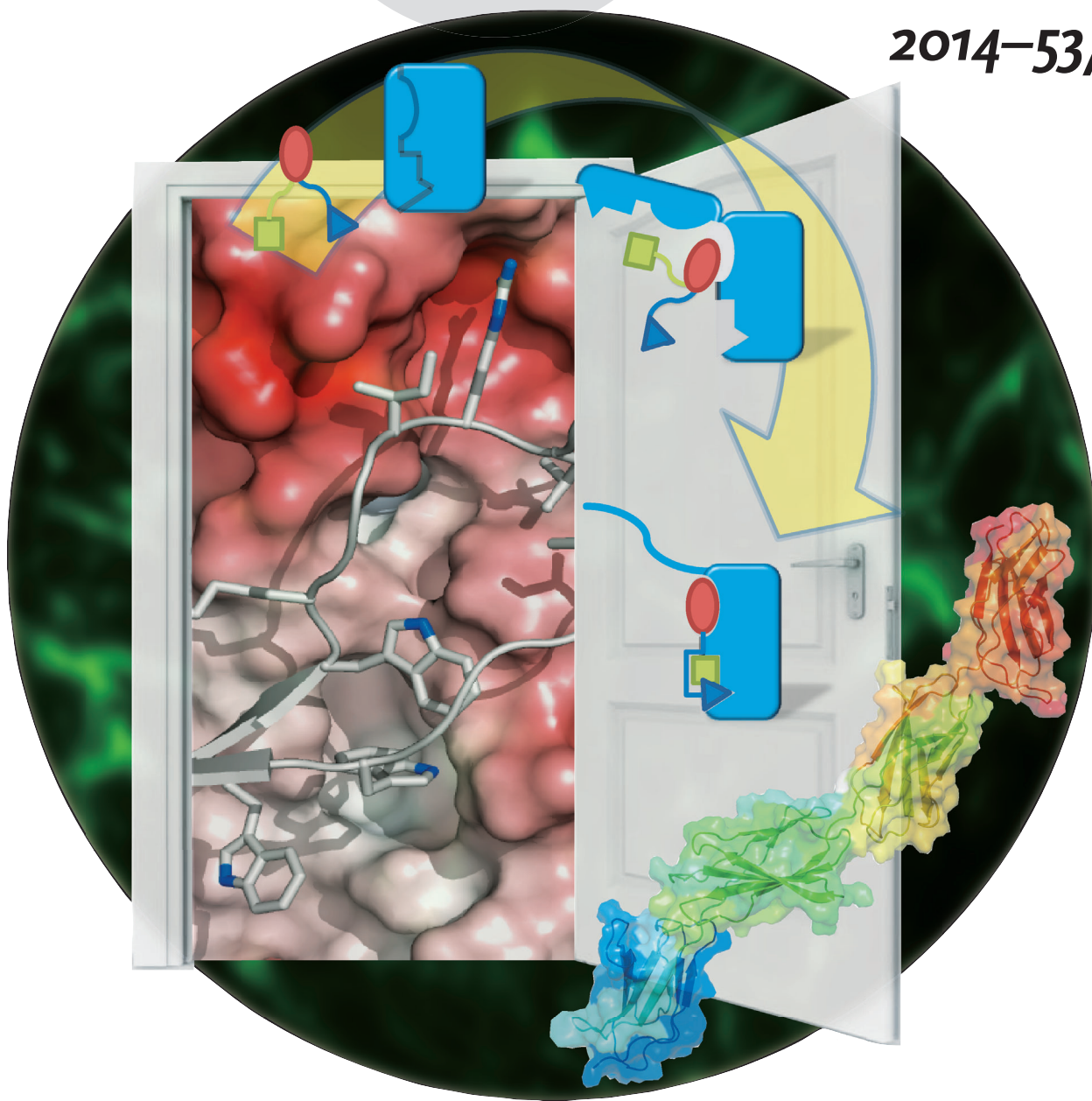
A Journal of the Gesellschaft Deutscher Chemiker

# Angewandte Chemie

GDCh  
International Edition

[www.angewandte.org](http://www.angewandte.org)

2014–53/37



## Protein–protein interactions ...

... can involve the folding of a disordered region to form the binding interface. In their Communication on page 9784 ff., J. Y. Suh and co-workers report the opposite case, in which binding is accompanied by local unfolding. The structure of an engineered peptide bound to fibronectin extradomain B reveals coupled unfolding and binding through  $\beta$ -strand displacement. The unfolding exposes a hydrophobic surface that provides key interactions for the complex.

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