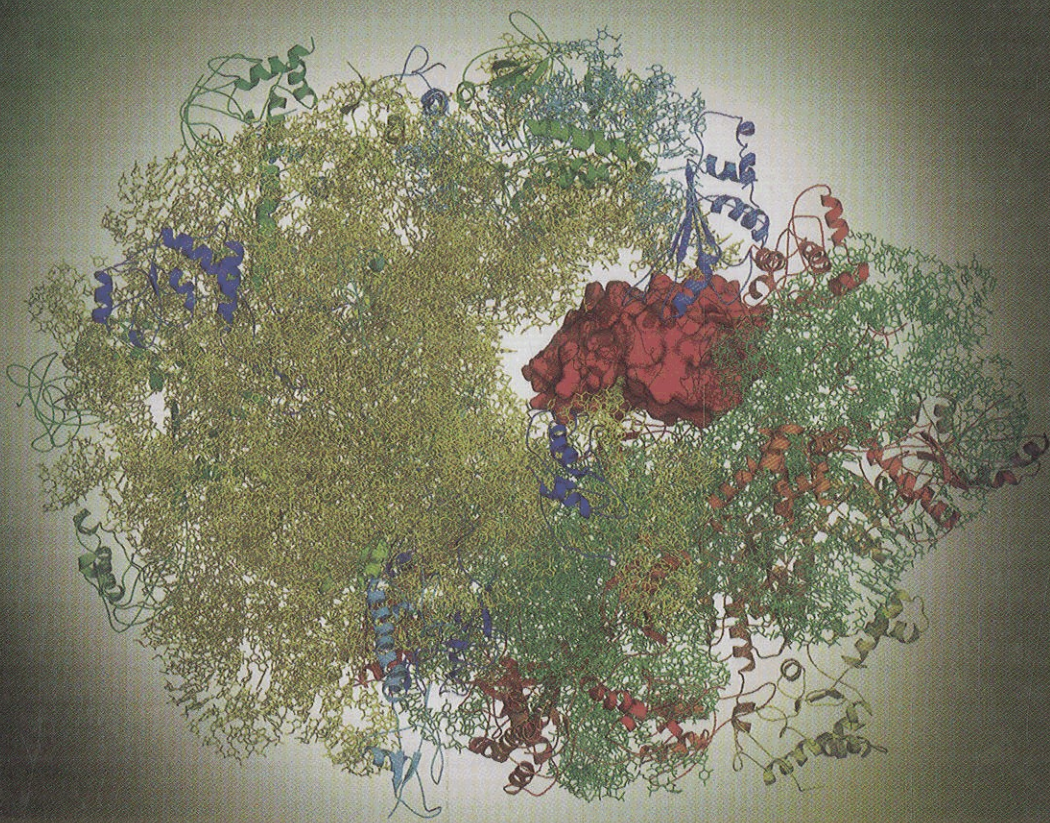




Prediction of Protein Structures, Functions and Interactions

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Contents

<i>List of Contributors</i>	ix
<i>Preface</i>	xiii
1 The Basics of Protein Sequence Analysis	1
<i>Katarzyna H. Kaminska, Kaja Milanowska and Janusz M. Bujnicki</i>	
2 First Steps of Protein Structure Prediction	39
<i>Karolina Majorek, Łukasz Kozłowski, Marcin Jąkałski and Janusz M. Bujnicki</i>	
3 Automated Prediction of Protein Function from Sequence	63
<i>Meghana Chitale, Troy Hawkins and Daisuke Kihara</i>	
4 Template Based Prediction of Three-dimensional Protein Structures: Fold Recognition and Comparative Modeling	87
<i>Jan Kosiński, Karolina L. Tkaczuk, Joanna M. Kasprzak and Janusz M. Bujnicki</i>	
5 Template-free Predictions of Three-dimensional Protein Structures: From First Principles to Knowledge-based Potentials	117
<i>Dominik Gront, Dorota Latek, Mateusz Kurcinski and Andrzej Kolinski</i>	
6 Quality Assessment of Protein Models	143
<i>Björn Wallner and Arne Elofsson</i>	
7 Prediction of Molecular Interactions from 3D-structures: From Small Ligands to Large Protein Complexes	159
<i>Kengo Kinoshita, Hidetoshi Kono and Kei Yura</i>	
8 Structure-based Prediction of Enzymes and Their Active Sites	187
<i>James W. Torrance and Janet M. Thornton</i>	
9 The Prediction of Macromolecular Complexes by Docking	211
<i>Sjoerd J. de Vries, Marc van Dijk and Alexandre M.J.J. Bonvin</i>	

10	Protein Function Prediction via Analysis of Interactomes	231
	<i>Elena Nabieva and Mona Singh</i>	
11	Integrating Prediction of Structure, Function, and Interactions	259
	<i>Michael Tress, Janusz M. Bujnicki, Gonzalo Lopez and Alfonso Valencia</i>	
	<i>Index</i>	281