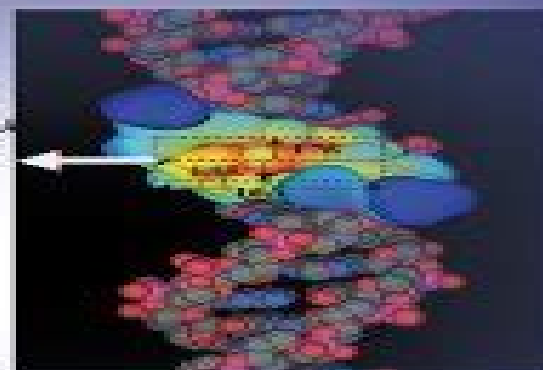
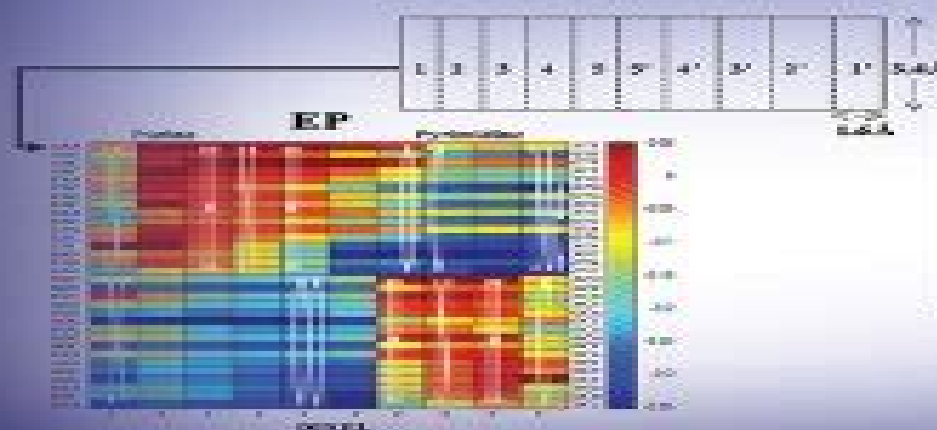


Computational Approaches in Cheminformatics and Bioinformatics

*Edited by
Rajarshi Guba and Andreas Bender*



Computational Approaches In Cheminformatics And Bioinformatics

**Navneet Sharma, Himanshu
Ojha, Pawan Raghav, Ramesh K. Goyal**



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terms of the number of new molecular entities NMEs Amongst the reasons noted for this are the adverse side effects and reduced efficiency of many of the potential compounds This is a problem both for the pharmaceutical industry and for those suffering from diseases for which there are no or few available treatments Advances in computational chemistry computer science structural biology and molecular biology have all contributed to improved drug design strategies and reduced the time taken for drug discovery By interfacing cheminformatics and bioinformatics with systems biology we can create a powerful tool for understanding the mechanisms of patho physiological systems and identifying lead molecules for various diseases This integration of drug design approaches can also play a crucial role in the prediction and rationalization of drug effects and side effects improving safety and efficacy and leading to better approval rates Addressing the lack of knowledge on the fundamental aspects of the various computational tools for drug discovery this book is a compilation of recent bioinformatics and cheminformatics approaches and their integration with systems biology Written primarily for researchers and academics in chem and bioinformatics it may also be a useful resource for advanced level students

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