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Features. Provides an overview of the basic principles of protein-protein interactions and the biophysical forces driving them; Investigates the architecture of

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Computational Protein-protein Interactions Hardcover. Often considered the workhorse of the cellular machinery, proteins are responsible for functions ranging from

Computational prediction of protein protein

Abstract: Protein interactions play an important role in the discovery of protein functions and pathways in biological processes. This is especially true in case of

Computational drug design targeting protein-

Computational Drug Design Targeting Protein-Protein Interactions. This review summarizes some current methods available for computational protein-protein

Computational investigation of senp:sumo protein-

How to Cite. Kumar, A. and Zhang, K. Y. J. (2013), Computational Investigation of SENP:SUMO Protein-Protein Interaction for Structure Based Drug Design.

Computational analysis of protein- protein

Computational analysis of protein-protein interactions: From sequences to networks. 28 September 3 October 2015 | Norwich, UK

Protein protein interaction prediction -

Protein protein interaction prediction is a field combining bioinformatics and structural biology in an attempt to identify and catalog physical interactions

Protein- protein interactions and networks -

The biological interactions of living organisms, and protein-protein interactions in particular, are astonishingly diverse and present numerous challenges

Protein- protein interactions | intechopen

Proteins are significant part of all biological events. The book provides concepts on Protein-Protein Interactions and methods on protein interactions. |Intechopen

Computational design of protein- protein

Computational design of protein protein interactions Tanja Kortemme^{1,2} and David Baker¹, Computational protein design strategies have been developed to reengineer

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View This Abstract Online; Computational drug design targeting protein-protein interactions. *Curr Pharm Des.* 2012; 18(9):1240-54 (ISSN: 1873-4286)

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Plos computational biology: deciphering protein

Citation: Shoemaker BA, Panchenko AR (2007) Deciphering Protein Protein Interactions. Part I. Experimental Techniques and Databases. *PLoS Comput Biol* 3(3): e42.

Protein protein interaction - wikipedia, the

Protein protein interactions (PPIs) refer to intentional physical contacts established between two or more proteins as a result of biochemical events and/or

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Computational methods of analysis of protein

computational and experimental methods for determining protein interactions, methods for validating interactions and ways to interpret the protein networks that

Computational design of protein protein

Abstract. Computational protein design strategies have been developed to reengineer protein protein interfaces in an automated, generalizable fashion.

Protein- protein interactions conference

Cambridge Healthtech Institute's conference on Protein-Protein Interactions has established itself as a successful and well attended event year after year. Now in

Computational identification of protein- protein

Plant protein-protein interaction networks have not been identified by large-scale experiments. In order to better understand the protein interactions in rice,

Plos computational biology: chapter 4: protein

Abstract. Proteins do not function in isolation; it is their interactions with one another and also with other molecules (e.g. DNA, RNA) that mediate metabolic and

Computational prediction of host-pathogen protein

Title: Computational prediction of host-pathogen protein-protein interactions. Publication Type: Journal Article: Year of Publication: 2007: Authors: Dyer, M. D., T

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FREE shipping on qualifying offers. Often considered the workhorse of

" computational study of protein- protein

Protein-protein interactions (PPIs) play important roles in biological systems. In particular, intra-protein interactions help create and maintain correctly folded

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Computational Prediction of Protein-Protein Interactions in Leishmania Predicted Proteomes Antonio M. Rezende 1,2 *, Edson L. Folador 1,3, Daniela de M. Resende 1,4

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