

# Ambrish Roy

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8012960/publications.pdf>

Version: 2024-02-01

8  
papers

11,277  
citations

1478280

6  
h-index

1588896

8  
g-index

9  
all docs

9  
docs citations

9  
times ranked

21107  
citing authors

#	ARTICLE	IF	CITATIONS
1	I-TASSER: a unified platform for automated protein structure and function prediction. Nature Protocols, 2010, 5, 725-738.	5.5	5,594
2	The I-TASSER Suite: protein structure and function prediction. Nature Methods, 2015, 12, 7-8.	9.0	4,923
3	COFACTOR: an accurate comparative algorithm for structure-based protein function annotation. Nucleic Acids Research, 2012, 40, W471-W477.	6.5	582
4	Recognizing Protein-Ligand Binding Sites by Global Structural Alignment and Local Geometry Refinement. Structure, 2012, 20, 987-997.	1.6	101
5	A Protocol for Computer-Based Protein Structure and Function Prediction. Journal of Visualized Experiments, 2011, , e3259.	0.2	54
6	QAUST: Protein Function Prediction Using Structure Similarity, Protein Interaction, and Functional Motifs. Genomics, Proteomics and Bioinformatics, 2021, 19, 998-1011.	3.0	14
7	Molecular and Structural Basis of Drift in the Functions of Closely-Related Homologous Enzyme Domains: Implications for Function Annotation Based on Homology Searches and Structural Genomics. In Silico Biology, 2009, 9, S41-S55.	0.4	6
8	Molecular and structural basis of drift in the functions of closely-related homologous enzyme domains: implications for function annotation based on homology searches and structural genomics. In Silico Biology, 2009, 9, S41-55.	0.4	2