

# Hailong Zhu

## List of Publications by Year in descending order

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1684188  
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times ranked

379  
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#	ARTICLE	IF	CITATIONS
1	Analysing the Effect of Mutation on Protein Function and Discovering Potential Inhibitors of CDK4: Molecular Modelling and Dynamics Studies. PLoS ONE, 2015, 10, e0133969.	2.5	50
2	Evolution- and Structure-Based Computational Strategy Reveals the Impact of Deleterious Missense Mutations on MODY 2 (Maturity-Onset Diabetes of the Young, Type 2). Theranostics, 2014, 4, 366-385.	10.0	48
3	Extrapolating the effect of deleterious nsSNPs in the binding adaptability of flavopiridol with CDK7 protein: a molecular dynamics approach. Human Genomics, 2013, 7, 10.	2.9	47
4	Integrating <i>In Silico</i> Prediction Methods, Molecular Docking, and Molecular Dynamics Simulation to Predict the Impact of ALK Missense Mutations in Structural Perspective. BioMed Research International, 2014, 2014, 1-14.	1.9	40
5	Evaluating Protein-protein Interaction (PPI) Networks for Diseases Pathway, Target Discovery, and Drug-design Using <i>In silico</i> Pharmacology™. Current Protein and Peptide Science, 2014, 15, 561-571.	1.4	19
6	CoagVDb: a comprehensive database for coagulation factors and their associated SAPs. Biological Research, 2015, 48, 35.	3.4	4
7	Rising Strengths Hong Kong SAR in Bioinformatics. Interdisciplinary Sciences, Computational Life Sciences, 2017, 9, 224-236.	3.6	3