Hailin Chen

List of Publications by Year in descending order

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		1040056	1199594	
13	378	9	12	
papers	citations	h-index	g-index	
13	13	13	493	
all docs	docs citations	times ranked	citing authors	

#	Article	IF	CITATIONS
1	A Semi-Supervised Method for Drug-Target Interaction Prediction with Consistency in Networks. PLoS ONE, 2013, 8, e62975.	2.5	98
2	Similarity-based methods for potential human microRNA-disease association prediction. BMC Medical Genomics, 2013, 6, 12.	1.5	95
3	Network-Based Inference Methods for Drug Repositioning. Computational and Mathematical Methods in Medicine, 2015, 2015, 1-7.	1.3	63
4	Prediction of Associations between OMIM Diseases and MicroRNAs by Random Walk on OMIM Disease Similarity Network. Scientific World Journal, The, 2013, 2013, 1-6.	2.1	30
5	Prediction and interpretation of miRNA-disease associations based on miRNA target genes using canonical correlation analysis. BMC Bioinformatics, 2019, 20, 404.	2.6	21
6	Prediction of Drug-Disease Associations for Drug Repositioning Through Drug-miRNA-Disease Heterogeneous Network. IEEE Access, 2018, 6, 45281-45287.	4.2	16
7	miRDDCR: a miRNA-based method to comprehensively infer drug-disease causal relationships. Scientific Reports, 2017, 7, 15921.	3.3	14
8	Comparative analysis of similarity measurements in miRNAs with applications to miRNA-disease association predictions. BMC Bioinformatics, 2020, 21, 176.	2.6	14
9	A miRNA-Driven Inference Model to Construct Potential Drug-Disease Associations for Drug Repositioning. BioMed Research International, 2015, 2015, 1-9.	1.9	12
10	In silico drug repositioning based on the integration of chemical, genomic and pharmacological spaces. BMC Bioinformatics, 2021, 22, 52.	2.6	6
11	Relating Disease-Gene Interaction Network With Disease-Associated ncRNAs. IEEE Access, 2019, 7, 133521-133528.	4.2	5
12	A Refined and Heuristic Algorithm for LD tagSNPs Selection. , 2011, , .		2
13	In silico drug repositioning based on integrated drug targets and canonical correlation analysis. BMC Medical Genomics, 2022, 15, 48.	1.5	2