

Yin Wang

List of Publications by Year in descending order

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12
papers

77
citations

1937685
4
h-index

1474206
9
g-index

13
all docs

13
docs citations

13
times ranked

121
citing authors

#	ARTICLE	IF	CITATIONS
1	Comparative Analysis of Multiple Neurodegenerative Diseases Based on Advanced Epigenetic Aging Brain. <i>Frontiers in Genetics</i> , 2021, 12, 657636.	2.3	6
2	The self-organization model reveals systematic characteristics of aging. <i>Theoretical Biology and Medical Modelling</i> , 2020, 17, 4.	2.1	1
3	The accelerated aging model reveals critical mechanisms of late-onset Parkinson's disease. <i>BioData Mining</i> , 2020, 13, 4.	4.0	6
4	The Model of Aging Acceleration Network Reveals the Correlation of Alzheimer's Disease and Aging at System Level. <i>BioMed Research International</i> , 2019, 2019, 1-11.	1.9	15
5	Network analysis of aging acceleration reveals systematic properties of 11 types of cancers. <i>FEBS Open Bio</i> , 2019, 9, 1292-1304.	2.3	2
6	Integrative Analysis of Dysfunctional Modules Driven by Genomic Alterations at System Level Across 11 Cancer Types. <i>Combinatorial Chemistry and High Throughput Screening</i> , 2019, 21, 771-783.	1.1	4
7	System modeling reveals the molecular mechanisms of HSC cell cycle alteration mediated by Mafk and Egr3 under leukemia. <i>BMC Systems Biology</i> , 2017, 11, 91.	3.0	9
8	Comprehensive analysis of differential co-expression patterns reveal transcriptional dysregulation mechanism and identify novel prognostic lncRNAs in esophageal squamous cell carcinoma. <i>OncoTargets and Therapy</i> , 2017, Volume 10, 3095-3105.	2.0	22
9	Motif-Based Text Mining of Microbial Metagenome Redundancy Profiling Data for Disease Classification. <i>BioMed Research International</i> , 2016, 2016, 1-11.	1.9	4
10	Integrative analysis of methylation and transcriptional profiles to predict aging and construct aging specific cross-tissue networks. <i>BMC Systems Biology</i> , 2016, 10, 132.	3.0	3
11	Quantitative Dynamic Modelling of the Gene Regulatory Network Controlling Adipogenesis. <i>PLoS ONE</i> , 2014, 9, e110563.	2.5	2
12	An improved dimensionality reduction method for meta-transcriptome indexing based diseases classification. <i>BMC Systems Biology</i> , 2012, 6, S12.	3.0	3