

Zeyneb Kurt

List of Publications by Year in descending order

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Version: 2024-02-01

22
papers

760
citations

1040056

9
h-index

1125743

13
g-index

28
all docs

28
docs citations

28
times ranked

1407
citing authors

#	ARTICLE	IF	CITATIONS
1	Transcription Factor MAFF (MAF Basic Leucine Zipper Transcription Factor F) Regulates an Atherosclerosis Relevant Network Connecting Inflammation and Cholesterol Metabolism. <i>Circulation</i> , 2021, 143, 1809-1823.	1.6	28
2	Integration of Multi-omics Data from Mouse Diversity Panel Highlights Mitochondrial Dysfunction in Non-alcoholic Fatty Liver Disease. <i>Cell Systems</i> , 2018, 6, 103-115.e7.	6.2	124
3	Tissue-specific pathways and networks underlying sexual dimorphism in non-alcoholic fatty liver disease. <i>Biology of Sex Differences</i> , 2018, 9, 46.	4.1	65
4	The Memory of Environmental Chemical Exposure in <i>C.Âlegans</i> Is Dependent on the Jumonji Demethylases <i>jmjd-2</i> and <i>jmjd-3/utx-1</i> . <i>Cell Reports</i> , 2018, 23, 2392-2404.	6.4	53
5	The Genetic Architecture of Diet-Induced Hepatic Fibrosis in Mice. <i>Hepatology</i> , 2018, 68, 2182-2196.	7.3	51
6	Applications and Limitations of Mouse Models for Understanding Human Atherosclerosis. <i>Cell Metabolism</i> , 2017, 25, 248-261.	16.2	161
7	Investigation of association estimators in network inference algorithms on breast cancer proteomic data. , 2017, , .		0
8	Estimation of the proteomic cancer co-expression sub networks by using association estimators. <i>PLoS ONE</i> , 2017, 12, e0188016.	2.5	3
9	Shared genetic regulatory networks for cardiovascular disease and type 2 diabetes in multiple populations of diverse ethnicities in the United States. <i>PLoS Genetics</i> , 2017, 13, e1007040.	3.5	82
10	Comprehensive review of association estimators for the inference of gene networks. <i>Turkish Journal of Electrical Engineering and Computer Sciences</i> , 2016, 24, 695-718.	1.4	11
11	Mergeomics: multidimensional data integration to identify pathogenic perturbations to biological systems. <i>BMC Genomics</i> , 2016, 17, 874.	2.8	106
12	Structural Analysis of Treatment Cycles Representing Transitions between Nursing Organizational Units Inferred from Diabetes. <i>PLoS ONE</i> , 2015, 10, e0127152.	2.5	0
13	Netmes: Assessing Gene Network Inference Algorithms by Network-Based Measures. <i>Evolutionary Bioinformatics</i> , 2014, 10, EBO.S13481.	1.2	3
14	A comprehensive comparison of association estimators for gene network inference algorithms. <i>Bioinformatics</i> , 2014, 30, 2142-2149.	4.1	21
15	Impacts of the different spline orders on the B-spline association estimator. , 2013, , .		0
16	Comparison of several classification algorithms for gender recognition from face images. , 2012, , .		11
17	Comparison of feature extraction and feature selection approaches to decide whether a face image belongs to a male or a female. , 2011, , .		1
18	Linear Discriminant Analysis in Ottoman Alphabet Character Recognition. <i>Lecture Notes in Electrical Engineering</i> , 2009, , 601-607.	0.4	4

#	ARTICLE	IF	CITATIONS
19	A simultaneous localization and map building algorithm based on sequential Monte Carlo method. , 2008, , .		0
20	Ottoman Alphabet Character Recognition by LDA. , 2007, , .		0
21	A Database of Non-Manual Signs in Turkish Sign Language. , 2007, , .		26
22	Integration of Multi-omics Data from Mouse Diversity Panel Highlights Mitochondrial Dysfunction in Non-Alcoholic Fatty Liver Disease. SSRN Electronic Journal, 0, , .	0.4	2