

Pujan Joshi

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/12189815/publications.pdf>

Version: 2024-02-01

15
papers

329
citations

1307594

7
h-index

1281871

11
g-index

15
all docs

15
docs citations

15
times ranked

562
citing authors

#	ARTICLE	IF	CITATIONS
1	Chemotherapy-induced oral mucositis is associated with detrimental bacterial dysbiosis. <i>Microbiome</i> , 2019, 7, 66.	11.1	140
2	Genome-wide analysis of FoxO1 binding in hepatic chromatin: Potential involvement of FoxO1 in linking retinoid signaling to hepatic gluconeogenesis. <i>Nucleic Acids Research</i> , 2012, 40, 11499-11509.	14.5	65
3	<i>GABRA2</i> Alcohol Dependence Risk Allele is Associated with Reduced Expression of Chromosome 4p12 GABA _A Subunit Genes in Human Neural Cultures. <i>Alcoholism: Clinical and Experimental Research</i> , 2015, 39, 1654-1664.	2.4	44
4	Astaxanthin attenuates the increase in mitochondrial respiration during the activation of hepatic stellate cells. <i>Journal of Nutritional Biochemistry</i> , 2019, 71, 82-89.	4.2	26
5	A route-based pathway analysis framework integrating mutation information and gene expression data. <i>Methods</i> , 2017, 124, 3-12.	3.8	16
6	rPAC: Route based pathway analysis for cohorts of gene expression data sets. <i>Methods</i> , 2022, 198, 76-87.	3.8	10
7	BioTarget: A Computational Framework Identifying Cancer Type Specific Transcriptional Targets of Immune Response Pathways. <i>Scientific Reports</i> , 2019, 9, 9029.	3.3	7
8	The LG/J murine strain exhibits near-normal tendon biomechanical properties following a full-length central patellar tendon defect. <i>Connective Tissue Research</i> , 2016, 57, 496-506.	2.3	6
9	PBC: A Software Framework Facilitating Pattern-Based Clustering for Microarray Data Analysis. , 2009, , .		4
10	A software framework integrating gene expression patterns, binding site analysis and gene ontology to hypothesize gene regulation relationships. , 2013, , .		4
11	Deep pathway analysis incorporating mutation information and gene expression data. , 2016, , .		2
12	Genome-wide distribution of 5hmC in the dental pulp of mouse molars and incisors. <i>Journal of Biochemistry</i> , 2022, 171, 123-129.	1.7	2
13	Predicting the targets of IRF8 and NFATc1 during osteoclast differentiation using the machine learning method framework cTAP. <i>BMC Genomics</i> , 2022, 23, 14.	2.8	2
14	The chromatin accessibility landscape in the dental pulp of mouse molars and incisors. <i>Acta Biochimica Polonica</i> , 2022, , .	0.5	1
15	Genome-Wide Analysis for Identifying FOXO Protein-Binding Sites. <i>Methods in Molecular Biology</i> , 2019, 1890, 193-203.	0.9	0