

Manjunath Kustagi

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6328185/publications.pdf>

Version: 2024-02-01

14
papers

1,592
citations

1163117

8
h-index

1281871

11
g-index

16
all docs

16
docs citations

16
times ranked

3830
citing authors

#	ARTICLE	IF	CITATIONS
1	Reverse engineering cellular networks. Nature Protocols, 2006, 1, 662-671.	12.0	345
2	A human B cell interactome identifies MYB and FOXM1 as master regulators of proliferation in germinal centers. Molecular Systems Biology, 2010, 6, 377.	7.2	336
3	A single-cell survey of the human first-trimester placenta and decidua. Science Advances, 2018, 4, eaau4788.	10.3	282
4	Tubular cell and keratinocyte single-cell transcriptomics applied to lupus nephritis reveal type I IFN and fibrosis relevant pathways. Nature Immunology, 2019, 20, 915-927.	14.5	275
5	Single cell RNA sequencing to dissect the molecular heterogeneity in lupus nephritis. JCI Insight, 2017, 2, .	5.0	164
6	Gene expression analysis uncovers similarity and differences among Burkitt lymphoma subtypes. Blood, 2011, 117, 3596-3608.	1.4	128
7	Interrogation of a Context-Specific Transcription Factor Network Identifies Novel Regulators of Pluripotency. Stem Cells, 2015, 33, 367-377.	3.2	32
8	The Cyni framework for network inference in Cytoscape. Bioinformatics, 2015, 31, 1499-1501.	4.1	9
9	Mechanism and Role of SOX2 Repression in Seminoma: Relevance to Human Germline Specification. Stem Cell Reports, 2016, 6, 772-783.	4.8	8
10	Elucidating synergistic dependencies in lung adenocarcinoma by proteome-wide signaling-network analysis. PLoS ONE, 2019, 14, e0208646.	2.5	6
11	300 Insights from single-cell RNA sequencing of skin and kidney in lupus nephritis. , 2019, , .		0
12	Abstract 4240: Transcription factor networks that regulate pluripotency and lineage differentiation in adult human male germ cell tumors. , 2010, , .		0
13	Abstract 3410: Regulation of pluripotency and lineage differentiation in human male germ cell tumors. , 2011, , .		0
14	Abstract PR11: Dissecting signaling transduction network to infer master regulators of non small cell lung cancer. , 2012, , .		0