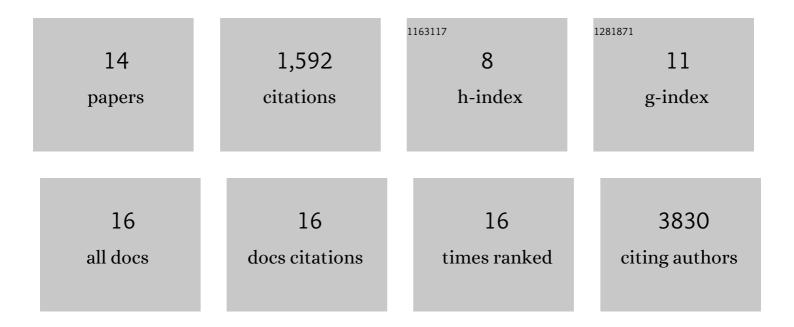
## Manjunath Kustagi

List of Publications by Year in descending order

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ΜΑΝΙΙΙΝΑΤΗ ΚΙΙSTACI

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