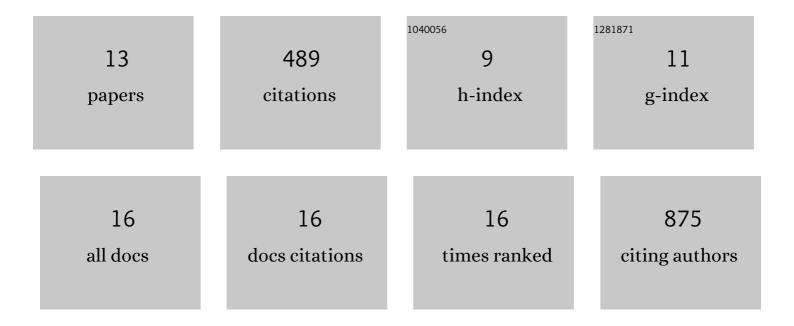
## Isha Sethi

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1420355/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Analysis of Cardiac Myocyte Maturation Using CASAAV, a Platform for Rapid Dissection of Cardiac Myocyte Gene Function In Vivo. Circulation Research, 2017, 120, 1874-1888.	4.5	106
2	A reference map of murine cardiac transcription factor chromatin occupancy identifies dynamic and conserved enhancers. Nature Communications, 2019, 10, 4907.	12.8	100
3	Hierarchical and stage-specific regulation of murine cardiomyocyte maturation by serum response factor. Nature Communications, 2018, 9, 3837.	12.8	63
4	A global analysis of the complex landscape of isoforms and regulatory networks of p63 in human cells and tissues. BMC Genomics, 2015, 16, 584.	2.8	52
5	Evolutionary re-wiring of p63 and the epigenomic regulatory landscape in keratinocytes and its potential implications on species-specific gene expression and phenotypes. Nucleic Acids Research, 2017, 45, 8208-8224.	14.5	39
6	Sarcomeres regulate murine cardiomyocyte maturation through MRTF-SRF signaling. Proceedings of the United States of America, 2021, 118, .	7.1	38
7	Role of chromatin and transcriptional co-regulators in mediating p63-genome interactions in keratinocytes. BMC Genomics, 2014, 15, 1042.	2.8	36
8	Massively parallel in vivo CRISPR screening identifies RNF20/40 as epigenetic regulators of cardiomyocyte maturation. Nature Communications, 2021, 12, 4442.	12.8	27
9	Reactivation of super-enhancers by KLF4 in human Head and Neck Squamous Cell Carcinoma. Oncogene, 2020, 39, 262-277.	5.9	15
10	Molecular Profiling Establishes Genetic Features Predictive of the Efficacy of the p110β Inhibitor KIN-193. Cancer Research, 2019, 79, 4524-4531.	0.9	7
11	De-Noising, Clustering, Classification, and Representation of Microarray Data for Disease Diagnostics. Advances in Computational Intelligence and Robotics Book Series, 2014, , 149-174.	0.4	0
12	Abstract B13: Genomic mistargeting of p63 drives the cancer phenotype in head and neck squamous cell carcinoma. , 2016, , .		0
13	Abstract A50: The identification of an Ets1-driven gene signature in head and neck squamous cell carcinoma. , 2016, , .		0