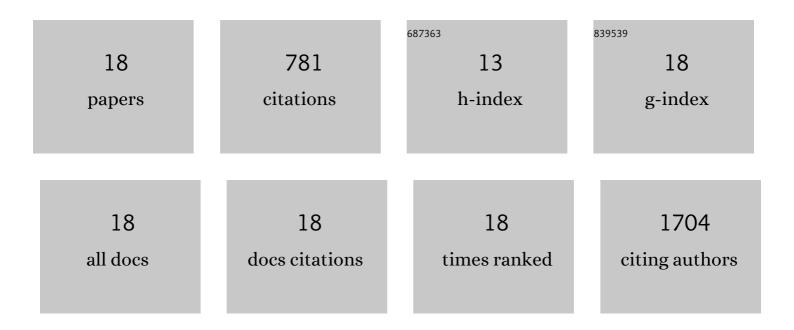
Muhammad Ahmad Maqbool

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	High-throughput and quantitative assessment of enhancer activity in mammals by CapStarr-seq. Nature Communications, 2015, 6, 6905.	12.8	138
2	Divergent transcription is associated with promoters of transcriptional regulators. BMC Genomics, 2013, 14, 914.	2.8	95
3	Differential Responses of Immune Cells to Type I Interferon Contribute to Host Resistance to Viral Infection. Cell Host and Microbe, 2012, 12, 571-584.	11.0	89
4	Tyrosine phosphorylation of RNA polymerase II CTD is associated with antisense promoter transcription and active enhancers in mammalian cells. ELife, 2014, 3, e02105.	6.0	76
5	Tyrosine-1 of RNA Polymerase II CTD Controls Global Termination of Gene Transcription in Mammals. Molecular Cell, 2018, 69, 48-61.e6.	9.7	66
6	ARS2 is a general suppressor of pervasive transcription. Nucleic Acids Research, 2017, 45, 10229-10241.	14.5	53
7	Site- and allele-specific polycomb dysregulation in T-cell leukaemia. Nature Communications, 2015, 6, 6094.	12.8	47
8	Dynamic recruitment of Ets1 to both nucleosome-occupied and -depleted enhancer regions mediates a transcriptional program switch during early T-cell differentiation. Nucleic Acids Research, 2016, 44, 3567-3585.	14.5	39
9	Regulation of the positive transcriptional effect of PLZF through a non-canonical EZH2 activity. Nucleic Acids Research, 2018, 46, 3339-3350.	14.5	26
10	A threshold level of NFATc1 activity facilitates thymocyte differentiation and opposes notch-driven leukaemia development. Nature Communications, 2016, 7, 11841.	12.8	23
11	Site-specific methylation and acetylation of lysine residues in the C-terminal domain (CTD) of RNA polymerase II. Transcription, 2015, 6, 91-101.	3.1	22
12	Pasha: a versatile R package for piling chromatin HTS data. Bioinformatics, 2016, 32, 2528-2530.	4.1	21
13	Regulation of Hepatitis C Virus Replication by Nuclear Translocation of Nonstructural 5A Protein and Transcriptional Activation of Host Genes. Journal of Virology, 2013, 87, 5523-5539.	3.4	18
14	AP-1 Signaling by Fra-1 Directly Regulates HMGA1 Oncogene Transcription in Triple-Negative Breast Cancers. Molecular Cancer Research, 2019, 17, 1999-2014.	3.4	15
15	Fra-1 regulates its target genes via binding to remote enhancers without exerting major control on chromatin architecture in triple negative breast cancers. Nucleic Acids Research, 2021, 49, 2488-2508.	14.5	15
16	MIR sequences recruit zinc finger protein ZNF768 to expressed genes. Nucleic Acids Research, 2019, 47, 700-715.	14.5	14
17	Alternative Enhancer Usage and Targeted Polycomb Marking Hallmark Promoter Choice during T Cell Differentiation. Cell Reports, 2020, 32, 108048.	6.4	13
18	An update on recent methods applied for deciphering the diversity of the noncoding RNA genome structure and function. Methods, 2013, 63, 3-17.	3.8	11