

Muhammad Ahmad Maqbool

List of Publications by Year in descending order

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Version: 2024-02-01

18
papers

781
citations

687363

13
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839539

18
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docs citations

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times ranked

1704
citing authors

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