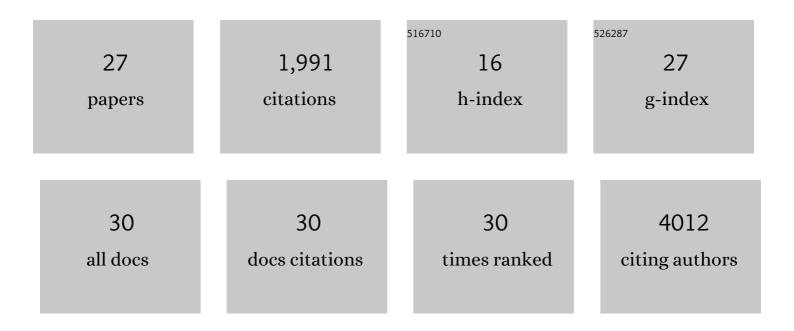
## Aditi Kanhere

List of Publications by Year in descending order

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Δριτι Κλνμερε

#	Article	IF	CITATIONS
1	Combined transcriptomic and phosphoproteomic analysis of BMP4 signaling in human embryonic stem cells. Stem Cell Research, 2021, 50, 102133.	0.7	3
2	Antisense RNAs during early vertebrate development are divided in groups with distinct features. Genome Research, 2021, 31, 995-1010.	5.5	7
3	A long intergenic non-coding RNA regulates nuclear localization of DNA methyl transferase-1. IScience, 2021, 24, 102273.	4.1	7
4	The Missing Link Between Cancer-Associated Variants and LncRNAs. Trends in Genetics, 2021, 37, 410-413.	6.7	4
5	8q24.21 Locus: A Paradigm to Link Non-Coding RNAs, Genome Polymorphisms and Cancer. International Journal of Molecular Sciences, 2021, 22, 1094.	4.1	15
6	Functional annotation of human long noncoding RNAs via molecular phenotyping. Genome Research, 2020, 30, 1060-1072.	5.5	109
7	Autoregulation of JARID2 through PRC2 interaction with its antisense ncRNA. BMC Research Notes, 2020, 13, 501.	1.4	0
8	A novel form of JARID2 is required for differentiation in lineage ommitted cells. EMBO Journal, 2019, 38, .	7.8	19
9	Regulation of Leukocytes by TspanC8 Tetraspanins and the "Molecular Scissor―ADAM10. Frontiers in Immunology, 2018, 9, 1451.	4.8	24
10	Exercise and high-fat feeding remodel transcript-metabolite interactive networks in mouse skeletal muscle. Scientific Reports, 2017, 7, 13485.	3.3	16
11	ldentification of RNA–Protein Interactions Through In Vitro RNA Pull-Down Assays. Methods in Molecular Biology, 2016, 1480, 99-113.	0.9	21
12	Exon junction complex proteins bind nascent transcripts independently of pre-mRNA splicing in Drosophila melanogaster. ELife, 2016, 5, .	6.0	19
13	Genome-Wide Regulatory Analysis Reveals That T-bet Controls Th17 Lineage Differentiation through Direct Suppression of IRF4. Journal of Immunology, 2013, 191, 5925-5932.	0.8	39
14	Modulation of Enhancer Looping and Differential Gene Targeting by Epstein-Barr Virus Transcription Factors Directs Cellular Reprogramming. PLoS Pathogens, 2013, 9, e1003636.	4.7	85
15	Genome-Wide Analyses of Zta Binding to the Epstein-Barr Virus Genome Reveals Interactions in both Early and Late Lytic Cycles and an Epigenetic Switch Leading to an Altered Binding Profile. Journal of Virology, 2012, 86, 12494-12502.	3.4	33
16	Downregulation of Integrin Receptor-Signaling Genes by Epstein-Barr Virus EBNA 3C via Promoter-Proximal and -Distal Binding Elements. Journal of Virology, 2012, 86, 5165-5178.	3.4	40
17	T-bet and GATA3 orchestrate Th1 and Th2 differentiation through lineage-specific targeting of distal regulatory elements. Nature Communications, 2012, 3, 1268.	12.8	292
18	Noncoding RNA localisation mechanisms in chromatin regulation. Silence: A Journal of RNA Regulation, 2012, 3, 2.	8.1	7

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#	Article	IF	CITATIONS
19	IL-2 Regulates Expression of <i>C-MAF</i> in Human CD4 T Cells. Journal of Immunology, 2011, 187, 3721-3729.	0.8	29
20	Short RNAs Are Transcribed from Repressed Polycomb Target Genes and Interact with Polycomb Repressive Complex-2. Molecular Cell, 2010, 38, 675-688.	9.7	338
21	Horizontal Gene Transfers in prokaryotes show differential preferences for metabolic and translational genes. BMC Evolutionary Biology, 2009, 9, 9.	3.2	44
22	CpG-depleted promoters harbor tissue-specific transcription factor binding signals—implications for motif overrepresentation analyses. Nucleic Acids Research, 2009, 37, 6305-6315.	14.5	44
23	Predicting transcription factor affinities to DNA from a biophysical model. Bioinformatics, 2007, 23, 134-141.	4.1	184
24	A novel method for prokaryotic promoter prediction based on DNA stability. BMC Bioinformatics, 2005, 6, 1.	2.6	462
25	Structural properties of promoters: similarities and differences between prokaryotes and eukaryotes. Nucleic Acids Research, 2005, 33, 3165-3175.	14.5	117
26	Identification of the domains for DNA binding and transactivation function of C protein from bacteriophage Mu. Proteins: Structure, Function and Bioinformatics, 2003, 52, 272-282.	2.6	6
27	An assessment of three dinucleotide parameters to predict DNA curvature by quantitative comparison with experimental data. Nucleic Acids Research, 2003, 31, 2647-2658.	14.5	22