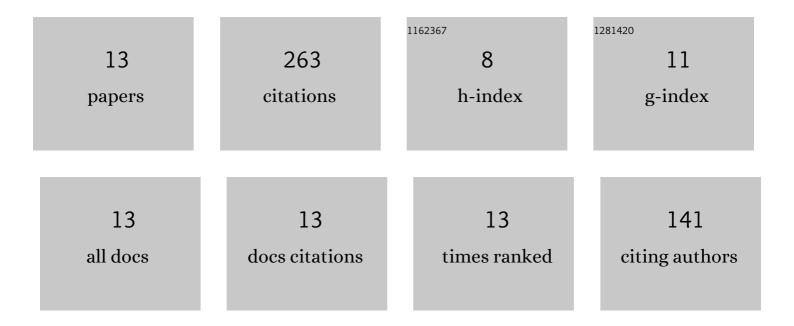
## Tasnim H Beacon

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6220155/publications.pdf Version: 2024-02-01



TASNIM H REACON

#	Article	IF	CITATIONS
1	The dynamic broad epigenetic (H3K4me3, H3K27ac) domain as a mark of essential genes. Clinical Epigenetics, 2021, 13, 138.	1.8	84
2	Epigenetic regulation of ACE2, the receptor of the SARS-CoV-2 virus <sup>1</sup> . Genome, 2021, 64, 386-399.	0.9	58
3	<scp>SARSâ€CoV</scp> â€2 multifaceted interaction with the human host. Part <scp>II</scp> : Innate immunity response, immunopathology, and epigenetics. IUBMB Life, 2020, 72, 2331-2354.	1.5	29
4	Genomic landscape of transcriptionally active histone arginine methylation marks, H3R2me2s and H4R3me2a, relative to nucleosome depleted regions. Gene, 2020, 742, 144593.	1.0	24
5	The chicken model organism for epigenomic research. Genome, 2021, 64, 476-489.	0.9	17
6	Atypical chromatin structure of immune-related genes expressed in chicken erythrocytes. Biochemistry and Cell Biology, 2020, 98, 171-177.	0.9	10
7	<scp>SARSâ€CoV</scp> â€2 multifaceted interaction with human host. Part I: What we have learnt and done so far, and the still unknown realities. IUBMB Life, 2020, 72, 2313-2330.	1.5	10
8	The key role of differential broad H3K4me3 and H3K4ac domains in breast cancer. Gene, 2022, 826, 146463.	1.0	9
9	Chromatin organization of transcribed genes in chicken polychromatic erythrocytes. Gene, 2019, 699, 80-87.	1.0	8
10	Transcriptionally Active Chromatin—Lessons Learned from the Chicken Erythrocyte Chromatin Fractionation. Cells, 2021, 10, 1354.	1.8	6
11	DNA methylation and chromatin modifications. , 2019, , 13-36.		4
12	Mitogen-induced transcriptional programming in human fibroblasts. Gene, 2021, 800, 145842.	1.0	4
13	The treatment of SARS-CoV2 with antivirals and mitigation of the cytokine storm syndrome: the role of gene expression. Genome, 2021, 64, 400-415.	0.9	0