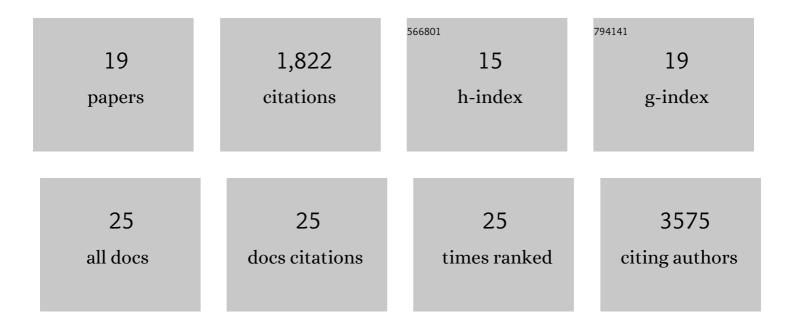
Namshik Han

List of Publications by Year in descending order

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ΝλΜςμικ Ηλν

#	Article	IF	CITATIONS
1	Promoter-bound METTL3 maintains myeloid leukaemia by m6A-dependent translation control. Nature, 2017, 552, 126-131.	13.7	833
2	Inflammatory Signals Induce AT2 Cell-Derived Damage-Associated Transient Progenitors that Mediate Alveolar Regeneration. Cell Stem Cell, 2020, 27, 366-382.e7.	5.2	303
3	The Forkhead Transcription Factor FOXM1 Controls Cell Cycle-Dependent Gene Expression through an Atypical Chromatin Binding Mechanism. Molecular and Cellular Biology, 2013, 33, 227-236.	1.1	185
4	Genomic positional conservation identifies topological anchor point RNAs linked to developmental loci. Genome Biology, 2018, 19, 32.	3.8	114
5	The forkhead transcription factor FOXK2 acts as a chromatin targeting factor for the BAP1-containing histone deubiquitinase complex. Nucleic Acids Research, 2014, 42, 6232-6242.	6.5	66
6	Ultradian Cortisol Pulsatility Encodes a Distinct, Biologically Important Signal. PLoS ONE, 2011, 6, e15766.	1.1	44
7	Long non-coding RNA ChRO1 facilitates ATRX/DAXX-dependent H3.3 deposition for transcription-associated heterochromatin reorganization. Nucleic Acids Research, 2018, 46, 11759-11775.	6.5	37
8	The FOXM1-PLK1 axis is commonly upregulated in oesophageal adenocarcinoma. British Journal of Cancer, 2012, 107, 1766-1775.	2.9	34
9	Identification of SARS-CoV-2–induced pathways reveals drug repurposing strategies. Science Advances, 2021, 7, .	4.7	34
10	<scp>DDX</scp> 3X <scp>RNA</scp> helicase affects breast cancer cell cycle progression by regulating expression of <scp>KLF</scp> 4. FEBS Letters, 2018, 592, 2308-2322.	1.3	32
11	Deregulation of the FOXM1 target gene network and its coregulatory partners in oesophageal adenocarcinoma. Molecular Cancer, 2015, 14, 69.	7.9	30
12	Current and prospective computational approaches and challenges for developing COVID-19 vaccines. Advanced Drug Delivery Reviews, 2021, 172, 249-274.	6.6	30
13	A novel long noncoding RNA Linc-ASEN represses cellular senescence through multileveled reduction of p21 expression. Cell Death and Differentiation, 2020, 27, 1844-1861.	5.0	23
14	Protein Kinase C Regulates Late Cell Cycle-Dependent Gene Expression. Molecular and Cellular Biology, 2012, 32, 4651-4661.	1.1	20
15	Methylation of histone H3 at lysine 37 by Set1 and Set2 prevents spurious DNA replication. Molecular Cell, 2021, 81, 2793-2807.e8.	4.5	18
16	Differential Expression of Soluble Receptor for Advanced Glycation End-products in Mice Susceptible or Resistant to Chronic Colitis. Inflammatory Bowel Diseases, 2020, 26, 360-368.	0.9	9
17	Progressive lung cancer determined by expression profiling and transcriptional regulation. International Journal of Oncology, 2012, 41, 242-52.	1.4	6
18	TIGERi: modeling and visualizing the responses to perturbation of a transcription factor network. BMC Bioinformatics, 2017, 18, 260.	1.2	2

#	Article	IF	CITATIONS
19	Identification of potential pan-coronavirus therapies using a computational drug repurposing platform. Methods, 2021, , .	1.9	1