

# Jin-Wu Nam

## List of Publications by Year in descending order

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Version: 2024-02-01

28  
papers

8,386  
citations

430874

18  
h-index

477307

29  
g-index

35  
all docs

35  
docs citations

35  
times ranked

16208  
citing authors

#	ARTICLE	IF	CITATIONS
1	Tumor immune microenvironment lncRNAs. <i>Briefings in Bioinformatics</i> , 2022, 23, .	6.5	77
2	hnRNPC induces isoform shifts in miR-21-5p leading to cancer development. <i>Experimental and Molecular Medicine</i> , 2022, 54, 812-824.	7.7	8
3	Preoperative immune landscape predisposes adverse outcomes in hepatocellular carcinoma patients with liver transplantation. <i>Npj Precision Oncology</i> , 2021, 5, 27.	5.4	11
4	The regulatory impact of RNA-binding proteins on microRNA targeting. <i>Nature Communications</i> , 2021, 12, 5057.	12.8	54
5	Single-cell transcriptome maps of myeloid blood cell lineages in <i>Drosophila</i> . <i>Nature Communications</i> , 2020, 11, 4483.	12.8	100
6	CGD: Comprehensive guide designer for CRISPR-Cas systems. <i>Computational and Structural Biotechnology Journal</i> , 2020, 18, 814-820.	4.1	6
7	A single-cell survey of <i>Drosophila</i> blood. <i>ELife</i> , 2020, 9, .	6.0	134
8	The small peptide world in long noncoding RNAs. <i>Briefings in Bioinformatics</i> , 2019, 20, 1853-1864.	6.5	183
9	UPF1/SMG7-dependent microRNA-mediated gene regulation. <i>Nature Communications</i> , 2019, 10, 4181.	12.8	20
10	<i>En bloc</i> and segmental deletions of human <i>XIST</i> reveal X chromosome inactivation-involving RNA elements. <i>Nucleic Acids Research</i> , 2019, 47, 3875-3887.	14.5	28
11	HERES, a lncRNA that regulates canonical and noncanonical Wnt signaling pathways via interaction with EZH2. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 24620-24629.	7.1	45
12	Comprehensive genome and transcriptome analyses reveal genetic relationship, selection signature, and transcriptome landscape of small-sized Korean native Jeju horse. <i>Scientific Reports</i> , 2019, 9, 16672.	3.3	18
13	HuR stabilizes a polyadenylated form of replication-independent histone mRNAs under stress conditions. <i>FASEB Journal</i> , 2019, 33, 2680-2693.	0.5	6
14	The present and future of <i>de novo</i> whole-genome assembly. <i>Briefings in Bioinformatics</i> , 2018, 19, bbw096.	6.5	139
15	TERIUS: accurate prediction of lncRNA via high-throughput sequencing data representing RNA-binding protein association. <i>BMC Bioinformatics</i> , 2018, 19, 41.	2.6	8
16	The long noncoding RNA LUCAT1 promotes tumorigenesis by controlling ubiquitination and stability of DNA methyltransferase 1 in esophageal squamous cell carcinoma. <i>Cancer Letters</i> , 2018, 417, 47-57.	7.2	112
17	Whole genome and transcriptome maps of the entirely black native Korean chicken breed Yeonsan Ogye. <i>GigaScience</i> , 2018, 7, .	6.4	20
18	Non-Coding Transcriptome Maps across Twenty Tissues of the Korean Black Chicken, Yeonsan Ogye. <i>International Journal of Molecular Sciences</i> , 2018, 19, 2359.	4.1	6

#	ARTICLE	IF	CITATIONS
19	High-confidence coding and noncoding transcriptome maps. <i>Genome Research</i> , 2017, 27, 1050-1062.	5.5	58
20	In vivo high-throughput profiling of CRISPR-Cpf1 activity. <i>Nature Methods</i> , 2017, 14, 153-159.	19.0	305
21	Incredible RNA: Dual Functions of Coding and Noncoding. <i>Molecules and Cells</i> , 2016, 39, 367-374.	2.6	89
22	Pseudo-Reference-Based Assembly of Vertebrate Transcriptomes. <i>Genes</i> , 2016, 7, 10.	2.4	2
23	Predicting effective microRNA target sites in mammalian mRNAs. <i>ELife</i> , 2015, 4, .	6.0	5,779
24	Global estimation of the 3' untranslated region landscape using RNA sequencing. <i>Methods</i> , 2015, 83, 111-117.	3.8	48
25	Global Analyses of the Effect of Different Cellular Contexts on MicroRNA Targeting. <i>Molecular Cell</i> , 2014, 53, 1031-1043.	9.7	276
26	Degradome sequencing reveals an endogenous microRNA target in <i>C. elegans</i> . <i>FEBS Letters</i> , 2013, 587, 964-969.	2.8	17
27	Long noncoding RNAs in <i>C. elegans</i> . <i>Genome Research</i> , 2012, 22, 2529-2540.	5.5	191
28	Expanding the MicroRNA Targeting Code: Functional Sites with Centered Pairing. <i>Molecular Cell</i> , 2010, 38, 789-802.	9.7	534