

# Anyou Wang

## List of Publications by Year in descending order

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Version: 2024-02-01

23  
papers

685  
citations

932766

10  
h-index

676716

22  
g-index

28  
all docs

28  
docs citations

28  
times ranked

1516  
citing authors

#	ARTICLE	IF	CITATIONS
1	Noncoding RNAs and Deep Learning Neural Network Discriminate Multi-Cancer Types. <i>Cancers</i> , 2022, 14, 352.	1.7	5
2	Noncoding RNAs endogenously rule the cancerous regulatory realm while proteins govern the normal. <i>Computational and Structural Biotechnology Journal</i> , 2022, 20, 1935-1945.	1.9	2
3	Distinctive functional regime of endogenous lncRNAs in dark regions of human genome. <i>Computational and Structural Biotechnology Journal</i> , 2022, 20, 2381-2390.	1.9	1
4	Noncoding RNAs Serve as the Deadliest Universal Regulators of all Cancers. <i>Cancer Genomics and Proteomics</i> , 2021, 18, 43-52.	1.0	11
5	The Impact of a Dried Fruit and Vegetable Supplement and Fiber Rich Shake on Gut and Health Parameters in Female Healthcare Workers: A Placebo-Controlled, Double-Blind, Randomized Clinical Trial. <i>Microorganisms</i> , 2021, 9, 843.	1.6	6
6	FINET: Fast Inferring NETwork. <i>BMC Research Notes</i> , 2020, 13, 521.	0.6	9
7	A combined biomarker and clinical panel for chronic graft versus host disease diagnosis. <i>Journal of Pathology: Clinical Research</i> , 2017, 3, 3-16.	1.3	23
8	Zika virus genome biology and molecular pathogenesis. <i>Emerging Microbes and Infections</i> , 2017, 6, 1-6.	3.0	99
9	Mining the human urine proteome for monitoring renal transplant injury. <i>Kidney International</i> , 2016, 89, 1244-1252.	2.6	80
10	Computational Models for Transplant Biomarker Discovery. <i>Frontiers in Immunology</i> , 2015, 6, 458.	2.2	11
11	Combined Targeted DNA Sequencing in Non-Small Cell Lung Cancer (NSCLC) Using UNCseq and NGScopy, and RNA Sequencing Using UNCqer for the Detection of Genetic Aberrations in NSCLC. <i>PLoS ONE</i> , 2015, 10, e0129280.	1.1	36
12	Systematically dissecting the global mechanism of miRNA functions in mouse pluripotent stem cells. <i>BMC Genomics</i> , 2015, 16, 490.	1.2	4
13	A Web-Server of Cell Type Discrimination System. <i>Scientific World Journal</i> , The, 2014, 2014, 1-5.	0.8	2
14	Alterations of LKB1 and KRAS and risk of brain metastasis: Comprehensive characterization by mutation analysis, copy number, and gene expression in non-small-cell lung carcinoma. <i>Lung Cancer</i> , 2014, 86, 255-261.	0.9	64
15	Transcription factor complex AP-1 mediates inflammation initiated by Chlamydia pneumoniae infection. <i>Cellular Microbiology</i> , 2013, 15, 779-794.	1.1	70
16	A Quantitative System for Discriminating Induced Pluripotent Stem Cells, Embryonic Stem Cells and Somatic Cells. <i>PLoS ONE</i> , 2013, 8, e56095.	1.1	10
17	Functional Modules Distinguish Human Induced Pluripotent Stem Cells from Embryonic Stem Cells. <i>Stem Cells and Development</i> , 2011, 20, 1937-1950.	1.1	26
18	A Systemic Network Triggered by Human Cytomegalovirus Entry. <i>Advances in Virology</i> , 2011, 2011, 1-11.	0.5	3

#	ARTICLE	IF	CITATIONS
19	A Systemic Network for <i>Chlamydia pneumoniae</i> Entry into Human Cells. Journal of Bacteriology, 2010, 192, 2809-2815.	1.0	21
20	The Arabidopsis A4 Subfamily of Lectin Receptor Kinases Negatively Regulates Abscisic Acid Response in Seed Germination. Plant Physiology, 2009, 149, 434-444.	2.3	69
21	Genome sequence divergences and functional variations in human cytomegalovirus strains. FEMS Immunology and Medical Microbiology, 2009, 55, 23-33.	2.7	10
22	Apoptosis, Cell Signaling, and Human Diseases: Molecular Mechanisms. JAMA - Journal of the American Medical Association, 2007, 298, 2200.	3.8	3
23	Global Gene Expression Responses to Cadmium Toxicity in Escherichia coli. Journal of Bacteriology, 2005, 187, 3259-3266.	1.0	112