Anyou Wang

List of Publications by Year
in descending order

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	Global Gene Expression Responses to Cadmium Toxicity in Escherichia coli. Journal of Bacteriology, 2005, 187, 3259-3266.	1.0	112
2	Zika virus genome biology and molecular pathogenesis. Emerging Microbes and Infections, 2017, 6, 1-6.	3.0	99
3	Mining the human urine proteome for monitoring renal transplant injury. Kidney International, 2016, 89, 1244-1252.	2.6	80
4	Transcription factor complex AP-1 mediates inflammation initiated by <i>Chlamydia pneumoniae</i> infection. Cellular Microbiology, 2013, 15, 779-794.	1.1	70
5	The Arabidopsis A4 Subfamily of Lectin Receptor Kinases Negatively Regulates Abscisic Acid Response in Seed Germination Â Â. Plant Physiology, 2009, 149, 434-444.	2.3	69
6	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. Lung Cancer, 2014, 86, 255-261.	0.9	64
7	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. PLoS ONE, 2015, 10, e0129280.	1.1	36
8	Functional Modules Distinguish Human Induced Pluripotent Stem Cells from Embryonic Stem Cells. Stem Cells and Development, 2011, 20, 1937-1950.	1.1	26
9	A combined biomarker and clinical panel for chronic graft versus host disease diagnosis. Journal of Pathology: Clinical Research, 2017, 3, 3-16.	1.3	23
10	A Systemic Network for <i>Chlamydia pneumoniae</i> Entry into Human Cells. Journal of Bacteriology, 2010, 192, 2809-2815.	1.0	21
11	Computational Models for Transplant Biomarker Discovery. Frontiers in Immunology, 2015, 6, 458.	2.2	11
12	Noncoding RNAs Serve as the Deadliest Universal Regulators of all Cancers. Cancer Genomics and Proteomics, 2021, 18, 43-52.	1.0	11
13	Genome sequence divergences and functional variations in human cytomegalovirus strains. FEMS Immunology and Medical Microbiology, 2009, 55, 23-33.	2.7	10
14	A Quantitative System for Discriminating Induced Pluripotent Stem Cells, Embryonic Stem Cells and Somatic Cells. PLoS ONE, 2013, 8, e56095.	1.1	10
15	FINET: Fast Inferring NETwork. BMC Research Notes, 2020, 13, 521.	0.6	9
16	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. Microorganisms, 2021, 9, 843.	1.6	6
17	Noncoding RNAs and Deep Learning Neural Network Discriminate Multi-Cancer Types. Cancers, 2022, 14, 352.	1.7	5
18	Systematically dissecting the global mechanism of miRNA functions in mouse pluripotent stem cells. BMC Genomics, 2015, 16, 490.	1.2	4

#	Article	IF	Citations
19	Apoptosis, Cell Signaling, and Human Diseases: Molecular Mechanisms. JAMA - Journal of the American Medical Association, 2007, 298, 2200.	3.8	3
20	A Systemic Network Triggered by Human Cytomegalovirus Entry. Advances in Virology, 2011, 2011, 1-11.	0.5	3
21	A Web-Server of Cell Type Discrimination System. Scientific World Journal, The, 2014, 2014, 1-5.	0.8	2
22	Noncoding RNAs endogenously rule the cancerous regulatory realm while proteins govern the normal. Computational and Structural Biotechnology Journal, 2022, 20, 1935-1945.	1.9	2
23	Distinctive functional regime of endogenous lncRNAs in dark regions of human genome. Computational and Structural Biotechnology Journal, 2022, 20, 2381-2390.	1.9	1