Dong Wang

List of Publications by Year in descending order

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Version: 2024-02-01

55 3,403 23 54
papers citations h-index g-index

56 56 56 6002 all docs citations times ranked citing authors

#	Article	IF	CITATIONS
1	Reprogramming transcription by distinct classes of enhancers functionally defined by eRNA. Nature, 2011, 474, 390-394.	13.7	777
2	The splicing factor SC35 has an active role in transcriptional elongation. Nature Structural and Molecular Biology, 2008, 15, 819-826.	3.6	316
3	Recurrently deregulated IncRNAs in hepatocellular carcinoma. Nature Communications, 2017, 8, 14421.	5.8	279
4	Acetylation-dependent regulation of PD-L1 nuclear translocation dictates the efficacy of anti-PD-1 immunotherapy. Nature Cell Biology, 2020, 22, 1064-1075.	4.6	182
5	Microarray is an efficient tool for circRNA profiling. Briefings in Bioinformatics, 2019, 20, 1420-1433.	3.2	161
6	Functional integration of transcriptional and RNA processing machineries. Current Opinion in Cell Biology, 2008, 20, 260-265.	2.6	154
7	Molecular basis for 5-carboxycytosine recognition by RNA polymerase II elongation complex. Nature, 2015, 523, 621-625.	13.7	141
8	Sensitive ChIP-DSL technology reveals an extensive estrogen receptor Â-binding program on human gene promoters. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 4852-4857.	3.3	120
9	Pre-mRNA splicing: where and when in the nucleus. Trends in Cell Biology, 2011, 21, 336-343.	3.6	118
10	An immune-related gene signature for predicting survival and immunotherapy efficacy in hepatocellular carcinoma. Cancer Immunology, Immunotherapy, 2021, 70, 967-979.	2.0	103
11	Nifuroxazide induces apoptosis and impairs pulmonary metastasis in breast cancer model. Cell Death and Disease, 2015, 6, e1701-e1701.	2.7	92
12	Network Pharmacology to Unveil the Biological Basis of Health-Strengthening Herbal Medicine in Cancer Treatment. Cancers, 2018, 10, 461.	1.7	83
13	An evolutionarily conserved DNA architecture determines target specificity of the TWIST family bHLH transcription factors. Genes and Development, 2015, 29, 603-616.	2.7	66
14	Versatile pathway-centric approach based on high-throughput sequencing to anticancer drug discovery. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 4609-4614.	3.3	63
15	Tissue-specific transcription reprogramming promotes liver metastasis of colorectal cancer. Cell Research, 2020, 30, 34-49.	5.7	60
16	TAp73-induced phosphofructokinase-1 transcription promotes the Warburg effect and enhances cell proliferation. Nature Communications, 2018, 9, 4683.	5.8	59
17	Noncoding RNAs Serve as Diagnosis and Prognosis Biomarkers for Hepatocellular Carcinoma. Clinical Chemistry, 2019, 65, 905-915.	1.5	57
18	Tumor immunological phenotype signature-based high-throughput screening for the discovery of combination immunotherapy compounds. Science Advances, 2021, 7, .	4.7	49

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19	Identification of high-confidence RNA regulatory elements by combinatorial classification of RNA–protein binding sites. Genome Biology, 2017, 18, 169.	3.8	42
20	Ribosome elongating footprints denoised by wavelet transform comprehensively characterize dynamic cellular translation events. Nucleic Acids Research, 2018, 46, e109-e109.	6.5	39
21	p15RS/RPRD1A (p15INK4b-related Sequence/Regulation of Nuclear Pre-mRNA Domain-containing Protein) Tj ETC Chemistry, 2015, 290, 9701-9713.	Qq1 1 0.7 1.6	84314 rgBT (34
22	Enhancing KDM5A and TLR activity improves the response to immune checkpoint blockade. Science Translational Medicine, 2020, 12, .	5.8	34
23	Super-Enhancer Redistribution as a Mechanism of Broad Gene Dysregulation in Repeatedly Drug-Treated Cancer Cells. Cell Reports, 2020, 31, 107532.	2.9	29
24	Guizhi Fuling Decoction inhibiting the PI3K and MAPK pathways in breast cancer cells revealed by HTS2 technology and systems pharmacology. Computational and Structural Biotechnology Journal, 2020, 18, 1121-1136.	1.9	27
25	Chemical genomics reveals inhibition of breast cancer lung metastasis by Ponatinib via c-Jun. Protein and Cell, 2019, 10, 161-177.	4.8	24
26	A large-scale transcriptional study reveals inhibition of COVID-19 related cytokine storm by traditional Chinese medicines. Science Bulletin, 2021, 66, 884-888.	4.3	24
27	Identification of a Tumor Immunological Phenotype-Related Gene Signature for Predicting Prognosis, Immunotherapy Efficacy, and Drug Candidates in Hepatocellular Carcinoma. Frontiers in Immunology, 2022, 13, 862527.	2.2	23
28	A comprehensive evaluation of connectivity methods for L1000 data. Briefings in Bioinformatics, 2020, 21, 2194-2205.	3.2	22
29	Comparison of Different Methods for Preparing Single Stranded DNA for Oligonucleotide Microarray. Analytical Letters, 2003, 36, 2849-2863.	1.0	20
30	Aureonitol Analogues and Orsellinic Acid Esters Isolated from <i>Chaetomium elatum</i> and Their Antineuroinflammatory Activity. Journal of Natural Products, 2021, 84, 3044-3054.	1.5	20
31	Evaluation of the immunomodulatory effects of anti-COVID-19 TCM formulae by multiple virus-related pathways. Signal Transduction and Targeted Therapy, 2021, 6, 50.	7.1	18
32	Histone H1 defect in escort cells triggers germline tumor in Drosophila ovary. Developmental Biology, 2017, 424, 40-49.	0.9	14
33	Single nucleotide polymorphism discrimination assisted by improved base stacking hybridization using oligonucleotide microarrays. BioTechniques, 2003, 35, 300-308.	0.8	12
34	The design and application of DNA chips for early detection of SARS-CoV from clinical samples. Journal of Clinical Virology, 2005, 33, 123-131.	1.6	11
35	Co-Expression of miR155 or LSD1 shRNA Increases the Anti-Tumor Functions of CD19 CAR-T Cells. Frontiers in Immunology, 2021, 12, 811364.	2.2	11
36	Cistrome plasticity and mechanisms of cistrome reprogramming. Cell Cycle, 2012, 11, 3199-3210.	1.3	10

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37	An automated microfluidic system for single-stranded DNA preparation and magnetic bead-based microarray analysis. Biomicrofluidics, 2015, 9, 024102.	1.2	10
38	Direct detection of 16S rRna using oligonucleotide microarrays assisted by base stacking hybridization and tyramide signal amplification. Journal of Proteomics, 2004, 59, 109-120.	2.4	9
39	Development of a base stacking hybridization-based microarray method for rapid identification of clinical isolates. Diagnostic Microbiology and Infectious Disease, 2007, 59, 149-156.	0.8	9
40	Functional Variants Identified Efficiently through an Integrated Transcriptome and Epigenome Analysis. Scientific Reports, 2018, 8, 2959.	1.6	9
41	Identification of Carpesium cernuum extract as a tumor migration inhibitor based on its biological response profiling in breast cancer cells. Phytomedicine, 2019, 64, 153072.	2.3	9
42	Dendrocalamus latiflorus and its component rutin exhibit glucose-lowering activities by inhibiting hepatic glucose production via AKT activation. Acta Pharmaceutica Sinica B, 2022, 12, 2239-2251.	5.7	8
43	Investigation of the Anti-Inflammatory Activity of Fusaproliferin Analogues Guided by Transcriptome Analysis. Frontiers in Pharmacology, 2022, 13, .	1.6	7
44	Oncogenic IncRNA LINCO0973 promotes Warburg effect by enhancing LDHA enzyme activity. Science Bulletin, 2021, 66, 1330-1341.	4.3	6
45	N-Acetyldopamine Dimer Attenuates DSS-Induced Ulcerative Colitis by Suppressing NF-κB and MAPK Pathways. Frontiers in Pharmacology, 2022, 13, 842730.	1.6	6
46	Two previously undescribed phthalides from <i>Talaromyces amestolkiae</i> , a symbiotic fungus of <i>Syngnathus acus</i> . Journal of Asian Natural Products Research, 2023, 25, 147-155.	0.7	6
47	ModuleRole: A Tool for Modulization, Role Determination and Visualization in Protein-Protein Interaction Networks. PLoS ONE, 2014, 9, e94608.	1.1	5
48	High-Throughput Strategies for the Discovery of Anticancer Drugs by Targeting Transcriptional Reprogramming. Frontiers in Oncology, 2021, 11, 762023.	1.3	5
49	Proliferatins suppress lipopolysaccharide-induced inflammation via inhibition of the NF-κB and MAPK signaling pathways. Bioorganic Chemistry, 2022, 124, 105810.	2.0	5
50	Detection of known mutations in hypertrophic cardiomyopathy using oligonucleotide microarrays assisted by improved base stacking hybridization. Biotechnology Letters, 2003, 25, 1613-1618.	1.1	4
51	Hydroanthraquinones from <i>Nigrospora sphaerica</i> and Their Anti-inflammatory Activity Uncovered by Transcriptome Analysis. Journal of Natural Products, 2022, 85, 1474-1485.	1.5	4
52	Taohong Siwu Decoction exerts anticancer effects on breast cancer via regulating MYC, BIRC5, EGF and PIK3R1 revealed by HTS2 technology. Computational and Structural Biotechnology Journal, 2022, 20, 3461-3472.	1.9	4
53	DNA interaction networks: an information highway for regulated gene expression in the 3-dimentional space of the nucleus. Cell Research, 2009, 19, 1316-1319.	5 . 7	1
54	IFNAR1 gene mutation may contribute to developmental stuttering in the Chinese population. Hereditas, 2021, 158, 46.	0.5	1

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55	A Novel Approach for Quality Control of Oligonucleotide Probes Using DHPLC. Analytical Letters, 2003, 36, 1463-1473.	1.0	O