

Huajing Teng

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

46
papers

660
citations

15
h-index

24
g-index

54
ext. papers

952
ext. citations

8.2
avg, IF

3.79
L-index

#	Paper	IF	Citations
46	Integrative analysis prioritised oxytocin-related biomarkers associated with the aetiology of autism spectrum disorder. <i>EBioMedicine</i> , 2022 , 81, 104091	8.8	0
45	Efficient correction of Duchenne muscular dystrophy mutations by SpCas9 and dual gRNAs. <i>Molecular Therapy - Nucleic Acids</i> , 2021 , 24, 403-415	10.7	5
44	OncoVar: an integrated database and analysis platform for oncogenic driver variants in cancers. <i>Nucleic Acids Research</i> , 2021 , 49, D1289-D1301	20.1	18
43	Population genomics reveal rapid genetic differentiation in a recently invasive population of <i>Rattus norvegicus</i> . <i>Frontiers in Zoology</i> , 2021 , 18, 6	2.8	1
42	Restricted Feeding Resets Endogenous Circadian Rhythm in Female Mice Under Constant Darkness. <i>Neuroscience Bulletin</i> , 2021 , 37, 1005-1009	4.3	1
41	Genomic and epigenomic evolution of acquired resistance to combination therapy in esophageal squamous cell carcinoma. <i>JCI Insight</i> , 2021 , 6,	9.9	1
40	Genomic analysis unveils mechanisms of northward invasion and signatures of plateau adaptation in the Asian house rat. <i>Molecular Ecology</i> , 2021 , 30, 6596-6610	5.7	3
39	Effect of Simultaneous Integrated Boost Intensity Modulated Radiation Therapy (SIB-IMRT) and Non-Operative Strategy on Outcomes of Distal Rectal Cancer Patients with Clinically Positive Lateral Pelvic Lymph Node. <i>Cancer Management and Research</i> , 2021 , 13, 537-546	3.6	0
38	Inter- and intratumor DNA methylation heterogeneity associated with lymph node metastasis and prognosis of esophageal squamous cell carcinoma. <i>Theranostics</i> , 2020 , 10, 3035-3048	12.1	10
37	Genetic evidence of gender difference in autism spectrum disorder supports the female-protective effect. <i>Translational Psychiatry</i> , 2020 , 10, 4	8.6	30
36	Prevalence and architecture of posttranscriptionally impaired synonymous mutations in 8,320 genomes across 22 cancer types. <i>Nucleic Acids Research</i> , 2020 , 48, 1192-1205	20.1	14
35	The clock-controlled chemokine contributes to neuroinflammation-induced depression. <i>FASEB Journal</i> , 2020 , 34, 8357-8366	0.9	4
34	Low intratumor heterogeneity correlates with increased response to PD-1 blockade in renal cell carcinoma. <i>Therapeutic Advances in Medical Oncology</i> , 2020 , 12, 1758835920977117	5.4	6
33	Co-expression Network of mRNAs and lncRNAs Regulated by Stress-Linked Behavioral Assays. <i>Psychopharmacology</i> , 2020 , 237, 571-582	4.7	6
32	Simultaneous Integrated Boost Intensity-Modulated Radiation Therapy Can Benefit the Locally Advanced Rectal Cancer Patients With Clinically Positive Lateral Pelvic Lymph Node. <i>Frontiers in Oncology</i> , 2020 , 10, 627572	5.3	1
31	Prediction of Clinical Outcome in Locally Advanced Non-Small Cell Lung Cancer Patients Treated With Chemoradiotherapy by Plasma Markers. <i>Frontiers in Oncology</i> , 2020 , 10, 625911	5.3	3
30	Whole-Exome Sequencing Identified a Novel Compound Heterozygous Genotype in a Chinese Han Patient with Argininosuccinate Lyase Deficiency. <i>BioMed Research International</i> , 2019 , 2019, 3530193	3	3

29	Transcriptome profiles of corticosterone-induced cytotoxicity reveals the involvement of neurite growth-related genes in depression. <i>Psychiatry Research</i> , 2019 , 276, 79-86	9.9	4
28	Androgen deprivation drives variation of androgen receptor trinucleotide repeats. <i>Acta Biochimica Et Biophysica Sinica</i> , 2019 , 51, 972-975	2.8	
27	OncoBase: a platform for decoding regulatory somatic mutations in human cancers. <i>Nucleic Acids Research</i> , 2019 , 47, D1044-D1055	20.1	18
26	VarCards: an integrated genetic and clinical database for coding variants in the human genome. <i>Nucleic Acids Research</i> , 2018 , 46, D1039-D1048	20.1	86
25	CirGRDB: a database for the genome-wide deciphering circadian genes and regulators. <i>Nucleic Acids Research</i> , 2018 , 46, D64-D70	20.1	19
24	EpiDenovo: a platform for linking regulatory de novo mutations to developmental epigenetics and diseases. <i>Nucleic Acids Research</i> , 2018 , 46, D92-D99	20.1	11
23	Altered expressions of memory genes in food-entrained circadian rhythm. <i>Acta Biochimica Et Biophysica Sinica</i> , 2018 , 50, 1068-1071	2.8	1
22	PAK2 Haploinsufficiency Results in Synaptic Cytoskeleton Impairment and Autism-Related Behavior. <i>Cell Reports</i> , 2018 , 24, 2029-2041	10.6	37
21	Transcriptomic signature associated with carcinogenesis and aggressiveness of papillary thyroid carcinoma. <i>Theranostics</i> , 2018 , 8, 4345-4358	12.1	41
20	Genetic landscape of papillary thyroid carcinoma in the Chinese population. <i>Journal of Pathology</i> , 2018 , 244, 215-226	9.4	56
19	Transcriptome Sequencing Reveals Candidate NF- κ B Target Genes Involved in Repeated Cocaine Administration. <i>International Journal of Neuropsychopharmacology</i> , 2018 , 21, 697-704	5.8	6
18	Asian house rats may facilitate their invasive success through suppressing brown rats in chronic interaction. <i>Frontiers in Zoology</i> , 2017 , 14, 20	2.8	9
17	Population Genomics Reveals Speciation and Introgression between Brown Norway Rats and Their Sibling Species. <i>Molecular Biology and Evolution</i> , 2017 , 34, 2214-2228	8.3	30
16	Dynamic Expression Changes in the Transcriptome of the Prefrontal Cortex after Repeated Exposure to Cocaine in Mice. <i>Frontiers in Pharmacology</i> , 2017 , 8, 142	5.6	11
15	Epigenetic Activation of in the Hippocampus Contributes to Depression-Like Behavior by Regulating D-Serine in Mice. <i>Frontiers in Molecular Neuroscience</i> , 2017 , 10, 139	6.1	12
14	RBP-Var: a database of functional variants involved in regulation mediated by RNA-binding proteins. <i>Nucleic Acids Research</i> , 2016 , 44, D154-63	20.1	31
13	Whole-Genome Sequencing Reveals Genetic Variation in the Asian House Rat. <i>G3: Genes, Genomes, Genetics</i> , 2016 , 6, 1969-77	3.2	6
12	Demethylation of c-MYB binding site mediates upregulation of Bdnf IV in cocaine-conditioned place preference. <i>Scientific Reports</i> , 2016 , 6, 22087	4.9	10

11	MBRidge: an accurate and cost-effective method for profiling DNA methylome at single-base resolution. <i>Journal of Molecular Cell Biology</i> , 2015 , 7, 299-313	6.3	4
10	Endogenous cellulolytic enzyme systems in the longhorn beetle <i>Mesosa myops</i> (Insecta: Coleoptera) studied by transcriptomic analysis. <i>Acta Biochimica Et Biophysica Sinica</i> , 2015 , 47, 741-8	2.8	13
9	Clock-controlled StAR α expression and corticosterone production contribute to the endotoxemia immune response. <i>Chronobiology International</i> , 2015 , 32, 358-67	3.6	8
8	Comparative RNA-seq analysis reveals potential mechanisms mediating the conversion to androgen independence in an LNCaP progression cell model. <i>Cancer Letters</i> , 2014 , 342, 130-8	9.9	16
7	Identification of a novel missense (C7W) mutation of SOD1 in a large familial amyotrophic lateral sclerosis pedigree. <i>Neurobiology of Aging</i> , 2014 , 35, 725.e11-5	5.6	6
6	Genome-wide identification and divergent transcriptional expression of StAR-related lipid transfer (START) genes in teleosts. <i>Gene</i> , 2013 , 519, 18-25	3.8	4
5	Regulation of peripheral clock to oscillation of substance P contributes to circadian inflammatory pain. <i>Anesthesiology</i> , 2012 , 117, 149-60	4.3	30
4	Deficiency of antinociception and excessive grooming induced by acute immobilization stress in <i>Per1</i> mutant mice. <i>PLoS ONE</i> , 2011 , 6, e16212	3.7	18
3	Evolutionary mode and functional divergence of vertebrate NMDA receptor subunit 2 genes. <i>PLoS ONE</i> , 2010 , 5, e13342	3.7	16
2	MagicViewer: integrated solution for next-generation sequencing data visualization and genetic variation detection and annotation. <i>Nucleic Acids Research</i> , 2010 , 38, W732-6	20.1	41
1	Expression profiling reveals a positive regulation by mPer2 on circadian rhythm of cytotoxicity receptors: <i>Ly49C</i> and <i>Nkg2d</i> . <i>Chronobiology International</i> , 2009 , 26, 1514-44	3.6	12