

Fan Wu

List of Publications by Citations

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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.

49
papers

1,183
citations

20
h-index

33
g-index

51
ext. papers

1,986
ext. citations

7.3
avg, IF

4.54
L-index

#	Paper	IF	Citations
49	mA RNA methylation regulators contribute to malignant progression and have clinical prognostic impact in gliomas. <i>Aging</i> , 2019 , 11, 1204-1225	5.6	125
48	Mutational Landscape of Secondary Glioblastoma Guides MET-Targeted Trial in Brain Tumor. <i>Cell</i> , 2018 , 175, 1665-1678.e18	56.2	125
47	Chinese Glioma Genome Atlas (CGGA): A Comprehensive Resource with Functional Genomic Data from Chinese Glioma Patients. <i>Genomics, Proteomics and Bioinformatics</i> , 2021 , 19, 1-12	6.5	103
46	An immune-related lncRNA signature for patients with anaplastic gliomas. <i>Journal of Neuro-Oncology</i> , 2018 , 136, 263-271	4.8	85
45	Molecular and clinical characterization of TIM-3 in glioma through 1,024 samples. <i>Oncotarget</i> , 2017 , 6, e1328339	7.2	74
44	Upregulation of long noncoding RNA HOXA-AS3 promotes tumor progression and predicts poor prognosis in glioma. <i>Oncotarget</i> , 2017 , 8, 53110-53123	3.3	44
43	LncRNA profile study reveals four-lncRNA signature associated with the prognosis of patients with anaplastic gliomas. <i>Oncotarget</i> , 2016 , 7, 77225-77236	3.3	44
42	Identification of an energy metabolism-related signature associated with clinical prognosis in diffuse glioma. <i>Aging</i> , 2018 , 10, 3185-3209	5.6	39
41	Prognostic power of a lipid metabolism gene panel for diffuse gliomas. <i>Journal of Cellular and Molecular Medicine</i> , 2019 , 23, 7741-7748	5.6	37
40	A three-gene signature for prognosis in patients with MGMT promoter-methylated glioblastoma. <i>Oncotarget</i> , 2016 , 7, 69991-69999	3.3	34
39	Bioinformatic analysis of gene expression and methylation regulation in glioblastoma. <i>Journal of Neuro-Oncology</i> , 2018 , 136, 495-503	4.8	30
38	Systematically characterize the clinical and biological significances of 1p19q genes in 1p/19q non-codeletion glioma. <i>Carcinogenesis</i> , 2019 , 40, 1229-1239	4.6	29
37	ALDH1A3 induces mesenchymal differentiation and serves as a predictor for survival in glioblastoma. <i>Cell Death and Disease</i> , 2018 , 9, 1190	9.8	27
36	Classification of diffuse lower-grade glioma based on immunological profiling. <i>Molecular Oncology</i> , 2020 , 14, 2081-2095	7.9	24
35	Acidosis enhances the self-renewal and mitochondrial respiration of stem cell-like glioma cells through CYP24A1-mediated reduction of vitamin D. <i>Cell Death and Disease</i> , 2019 , 10, 25	9.8	23
34	Molecular classification of IDH-mutant glioblastomas based on gene expression profiles. <i>Carcinogenesis</i> , 2019 , 40, 853-860	4.6	22
33	Expression profile analysis of antisense long non-coding RNA identifies WDFY3-AS2 as a prognostic biomarker in diffuse glioma. <i>Cancer Cell International</i> , 2018 , 18, 107	6.4	21

32	Combinations of four or more CpGs methylation present equivalent predictive value for MGMT expression and temozolomide therapeutic prognosis in gliomas. <i>CNS Neuroscience and Therapeutics</i> , 2019 , 25, 314-322	6.8	20
31	The antibiotic clofoctol suppresses glioma stem cell proliferation by activating KLF13. <i>Journal of Clinical Investigation</i> , 2019 , 129, 3072-3085	15.9	20
30	Amino acid metabolism-related gene expression-based risk signature can better predict overall survival for glioma. <i>Cancer Science</i> , 2019 , 110, 321-333	6.9	20
29	Elevated signature of a gene module coexpressed with CDC20 marks genomic instability in glioma. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019 , 116, 6975-6984	11.5	19
28	Molecular subtyping reveals immune alterations in IDH wild-type lower-grade diffuse glioma. <i>Journal of Pathology</i> , 2020 , 251, 272-283	9.4	19
27	A novel analytical model of MGMT methylation pyrosequencing offers improved predictive performance in patients with gliomas. <i>Modern Pathology</i> , 2019 , 32, 4-15	9.8	19
26	Long noncoding RNA HOXD-AS2 regulates cell cycle to promote glioma progression. <i>Journal of Cellular Biochemistry</i> , 2018 , 120, 8343	4.7	16
25	Surveying brain tumor heterogeneity by single-cell RNA-sequencing of multi-sector biopsies. <i>National Science Review</i> , 2020 , 7, 1306-1318	10.8	15
24	Systematically profiling the expression of eIF3 subunits in glioma reveals the expression of eIF3i has prognostic value in IDH-mutant lower grade glioma. <i>Cancer Cell International</i> , 2019 , 19, 155	6.4	14
23	Identification of an ATP metabolism-related signature associated with prognosis and immune microenvironment in gliomas. <i>Cancer Science</i> , 2020 , 111, 2325-2335	6.9	14
22	A Novel DNA Methylation-Based Signature Can Predict the Responses of MGMT Promoter Unmethylated Glioblastomas to Temozolomide. <i>Frontiers in Genetics</i> , 2019 , 10, 910	4.5	13
21	RNA processing genes characterize RNA splicing and further stratify lower-grade glioma. <i>JCI Insight</i> , 2019 , 5,	9.9	13
20	A three ion channel genes-based signature predicts prognosis of primary glioblastoma patients and reveals a chemotherapy sensitive subtype. <i>Oncotarget</i> , 2016 , 7, 74895-74903	3.3	11
19	Siglecs, Novel Immunotherapy Targets, Potentially Enhance The Effectiveness of Existing Immune Checkpoint Inhibitors in Glioma Immunotherapy. <i>OncoTargets and Therapy</i> , 2019 , 12, 10263-10273	4.4	11
18	Glioma-associated human endothelial cell-derived extracellular vesicles specifically promote the tumorigenicity of glioma stem cells via CD9. <i>Oncogene</i> , 2019 , 38, 6898-6912	9.2	10
17	Single-Cell RNA-Sequencing Shift in the Interaction Pattern Between Glioma Stem Cells and Immune Cells During Tumorigenesis. <i>Frontiers in Immunology</i> , 2020 , 11, 581209	8.4	10
16	Gene Expression Profiling Stratifies IDH-Wildtype Glioblastoma With Distinct Prognoses. <i>Frontiers in Oncology</i> , 2019 , 9, 1433	5.3	8
15	indicates assembly of M0 macrophage and more malignant phenotypes of glioma. <i>Aging</i> , 2020 , 12, 8397-8412	5.6	7

14	Predictive value of MGMT promoter methylation on the survival of TMZ treated -mutant glioblastoma. <i>Cancer Biology and Medicine</i> , 2021 , 18, 272-282	5.2	6
13	A comprehensive review of available omics data resources and molecular profiling for precision glioma studies. <i>Biomedical Reports</i> , 2019 , 10, 3-9	1.8	5
12	An MRI radiomics approach to predict survival and tumour-infiltrating macrophages in gliomas.. <i>Brain</i> , 2022 ,	11.2	5
11	Prognostic value of ion channel genes in Chinese patients with gliomas based on mRNA expression profiling. <i>Journal of Neuro-Oncology</i> , 2017 , 134, 397-405	4.8	4
10	Clinical Characterization and Immunosuppressive Regulation of CD161 (KLRB1) in Glioma through 916 Samples. <i>Cancer Science</i> , 2021 ,	6.9	4
9	A PTEN-COL17A1 fusion gene and its novel regulatory role in Collagen XVII expression and GBM malignance. <i>Oncotarget</i> , 2017 , 8, 85794-85803	3.3	4
8	Clinical and Biological Significances of a Methyltransferase-Related Signature in Diffuse Glioma. <i>Frontiers in Oncology</i> , 2020 , 10, 508	5.3	2
7	Characterization and prognostic significance of alternative splicing events in lower-grade diffuse gliomas. <i>Journal of Cellular and Molecular Medicine</i> , 2020 , 24, 13171-13180	5.6	2
6	Galectin-9/TIM-3 as a Key Regulator of Immune Response in Gliomas With Chromosome 1p/19q Codeletion.. <i>Frontiers in Immunology</i> , 2021 , 12, 800928	8.4	2
5	RPP30, a transcriptional regulator, is a potential pathogenic factor in glioblastoma. <i>Aging</i> , 2020 , 12, 16155616171	5.5	1
4	A novel DNA repair-related nomogram predicts survival in low-grade gliomas. <i>CNS Neuroscience and Therapeutics</i> , 2021 , 27, 186-195	6.8	1
3	Molecular Characterization and Clinical Relevance of in Gliomas 1,018 Chinese Cohort Patients.. <i>Frontiers in Cell and Developmental Biology</i> , 2021 , 9, 777182	5.7	0
2	Metabolic expression profiling stratifies diffuse lower-grade glioma into three distinct tumour subtypes. <i>British Journal of Cancer</i> , 2021 , 125, 255-264	8.7	0
1	Norepinephrine in Goal-Directed Fluid Therapy During General Anesthesia in Elderly Patients Undergoing Spinal Operation: Determining Effective Infusion Rate to Enhance Postoperative Functions.. <i>Current Genomics</i> , 2021 , 22, 620-629	2.6	