

Zhaoshi Bao

List of Publications by Year in Descending Order

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Version: 2024-04-26

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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.

31
papers

1,094
citations

16
h-index

33
g-index

33
ext. papers

1,671
ext. citations

6.8
avg, IF

3.96
L-index

| # | Paper | IF | Citations |
|----|---|------|-----------|
| 31 | Comprehensive analysis of the LncRNAs, MiRNAs, and MRNAs acting within the competing endogenous RNA network of LGG.. <i>Genetica</i> , 2022 , 150, 41 | 1.5 | 0 |
| 30 | Chemoradiotherapy with temozolomide vs. radiotherapy alone in patients with IDH wild-type and TERT promoter mutation WHO grade II/III gliomas: A prospective randomized study.. <i>Radiotherapy and Oncology</i> , 2021 , 167, 1-6 | 5.3 | 0 |
| 29 | 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 |
| 28 | Intratumor heterogeneity, microenvironment, and mechanisms of drug resistance in glioma recurrence and evolution. <i>Frontiers of Medicine</i> , 2021 , 15, 551-561 | 12 | 9 |
| 27 | Is Relating With M2 Macrophage and Remarkable Malignancy Characters in Low-Grade Glioma. <i>Frontiers in Immunology</i> , 2021 , 12, 659659 | 8.4 | 4 |
| 26 | Clinical practice guidelines for the management of adult diffuse gliomas. <i>Cancer Letters</i> , 2021 , 499, 60-73. | 9 | 61 |
| 25 | Comprehensive transcriptomic characterization reveals core genes and module associated with immunological changes via 1619 samples of brain glioma. <i>Cell Death and Disease</i> , 2021 , 12, 1140 | 9.8 | 2 |
| 24 | LINC00174 is a favorable prognostic biomarker in glioblastoma via promoting proliferative phenotype. <i>Cancer Biomarkers</i> , 2020 , 28, 421-427 | 3.8 | 6 |
| 23 | relevant bioinformatic profiling and prognostic value in gliomas. <i>Future Oncology</i> , 2020 , 16, 4279-4288 | 3.6 | 11 |
| 22 | MGMT genomic rearrangements contribute to chemotherapy resistance in gliomas. <i>Nature Communications</i> , 2020 , 11, 3883 | 17.4 | 47 |
| 21 | MEGF10, a Glioma Survival-Associated Molecular Signature, Predicts IDH Mutation Status. <i>Disease Markers</i> , 2018 , 2018, 5975216 | 3.2 | 8 |
| 20 | Identification of IDH-mutant gliomas by a prognostic signature according to gene expression profiling. <i>Aging</i> , 2018 , 10, 1977-1988 | 5.6 | 5 |
| 19 | NCMP-28. PTPRZ1-MET SIGNALING PROMOTES GLIOMA PROGRESSION THROUGH STIMULATION THE TRANSFORMATION FROM M1 TO M2 MACROPHAGE. <i>Neuro-Oncology</i> , 2018 , 20, vi199-vi199 | 1 | 1 |
| 18 | Mutational Landscape of Secondary Glioblastoma Guides MET-Targeted Trial in Brain Tumor. <i>Cell</i> , 2018 , 175, 1665-1678.e18 | 56.2 | 125 |
| 17 | Bioinformatic analyses reveal a distinct Notch activation induced by STAT3 phosphorylation in the mesenchymal subtype of glioblastoma. <i>Journal of Neurosurgery</i> , 2017 , 126, 249-259 | 3.2 | 15 |
| 16 | CDC20 with malignant progression and poor prognosis of astrocytoma revealed by analysis on gene expression. <i>Journal of Neuro-Oncology</i> , 2017 , 133, 87-95 | 4.8 | 13 |
| 15 | Interplay between PCBP2 and miRNA modulates ARHGDI1 expression and function in glioma migration and invasion. <i>Oncotarget</i> , 2016 , 7, 19483-98 | 3.3 | 32 |

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|----|---|------|-----|
| 14 | KIF23 is an independent prognostic biomarker in glioma, transcriptionally regulated by TCF-4. <i>Oncotarget</i> , 2016 , 7, 24646-55 | 3.3 | 25 |
| 13 | CGCG clinical practice guidelines for the management of adult diffuse gliomas. <i>Cancer Letters</i> , 2016 , 375, 263-273 | 9.9 | 253 |
| 12 | ALDH1A3: A Marker of Mesenchymal Phenotype in Gliomas Associated with Cell Invasion. <i>PLoS ONE</i> , 2015 , 10, e0142856 | 3.7 | 22 |
| 11 | PTBP1 induces ADAR1 p110 isoform expression through IRES-like dependent translation control and influences cell proliferation in gliomas. <i>Cellular and Molecular Life Sciences</i> , 2015 , 72, 4383-97 | 10.3 | 20 |
| 10 | Hypomethylated Rab27b is a progression-associated prognostic biomarker of glioma regulating MMP-9 to promote invasion. <i>Oncology Reports</i> , 2015 , 34, 1503-9 | 3.5 | 13 |
| 9 | Genetic and clinical characteristics of primary and secondary glioblastoma is associated with differential molecular subtype distribution. <i>Oncotarget</i> , 2015 , 6, 7318-24 | 3.3 | 33 |
| 8 | Integrated analysis using methylation and gene expression microarrays reveals PDE4C as a prognostic biomarker in human glioma. <i>Oncology Reports</i> , 2014 , 32, 250-60 | 3.5 | 10 |
| 7 | MicroRNA expression patterns in the malignant progression of gliomas and a 5-microRNA signature for prognosis. <i>Oncotarget</i> , 2014 , 5, 12908-15 | 3.3 | 48 |
| 6 | SOCS3 promoter hypermethylation is a favorable prognosticator and a novel indicator for G-CIMP-positive GBM patients. <i>PLoS ONE</i> , 2014 , 9, e91829 | 3.7 | 19 |
| 5 | Identification of miRNA-mediated core gene module for glioma patient prediction by integrating high-throughput miRNA, mRNA expression and pathway structure. <i>PLoS ONE</i> , 2014 , 9, e96908 | 3.7 | 23 |
| 4 | Epigenetic suppression of EGFR signaling in G-CIMP+ glioblastomas. <i>Oncotarget</i> , 2014 , 5, 7342-56 | 3.3 | 18 |
| 3 | BMP4, a strong better prognosis predictor, has a subtype preference and cell development association in gliomas. <i>Journal of Translational Medicine</i> , 2013 , 11, 100 | 8.5 | 27 |
| 2 | Molecular classification of gliomas based on whole genome gene expression: a systematic report of 225 samples from the Chinese Glioma Cooperative Group. <i>Neuro-Oncology</i> , 2012 , 14, 1432-40 | 1 | 133 |
| 1 | Chinese Glioma Genome Atlas (CGGA): A Comprehensive Resource with Functional Genomic Data for Chinese Glioma Patients | | 8 |