## Ke-Nan Zhang

List of Publications by Year in descending order

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430442 525886 1,416 29 18 27 citations h-index g-index papers 31 31 31 1342 docs citations times ranked citing authors all docs

| #  | Article  | IF  | CITATIONS |
|----|--|-----|-----------|
| 1  | Glioma-related epilepsy in patients with diffuse high-grade glioma after the 2016 WHO update: seizure characteristics, risk factors, and clinical outcomes. Journal of Neurosurgery, 2022, 136, 67-75.                             | 0.9 | 15        |
| 2  | Prognostic Pathways Guide Drug Indications in Pan-Cancers. Frontiers in Oncology, 2022, 12, 849552.  | 1.3 | 0         |
| 3  | <i>LRRFIP1</i> , an epigenetically regulated gene, is a prognostic biomarker and predicts malignant phenotypes of glioma. CNS Neuroscience and Therapeutics, 2022, 28, 873-883.  | 1.9 | 1         |
| 4  | Chinese Glioma Genome Atlas (CGGA): A Comprehensive Resource with Functional Genomic Data from Chinese Glioma Patients. Genomics, Proteomics and Bioinformatics, 2021, 19, 1-12.   | 3.0 | 439       |
| 5  | Plasminogen Activator Urokinase Receptor Implies Immunosuppressive Features and Acts as an Unfavorable Prognostic Biomarker in Glioma. Oncologist, 2021, 26, e1460-e1469.  | 1.9 | 21        |
| 6  | SAMD9 Is Relating With M2 Macrophage and Remarkable Malignancy Characters in Low-Grade Glioma. Frontiers in Immunology, 2021, 12, 659659.  | 2.2 | 16        |
| 7  | New-Onset Postoperative Seizures in Patients With Diffuse Gliomas: A Risk Assessment Analysis. Frontiers in Neurology, 2021, 12, 682535.   | 1.1 | 3         |
| 8  | METTL3 enhances the stability of MALAT1 with the assistance of HuR via m6A modification and activates NF-κB to promote the malignant progression of IDH-wildtype glioma. Cancer Letters, 2021, 511, 36-46.                         | 3.2 | 86        |
| 9  | YTHDF2 facilitates UBXN1 mRNA decay by recognizing METTL3-mediated m6A modification to activate NF- $^{\rm lp}$ B and promote the malignant progression of glioma. Journal of Hematology and Oncology, 2021, 14, 109.              | 6.9 | 92        |
| 10 | Molecular subtype impacts surgical resection in low-grade gliomas: A Chinese Glioma Genome Atlas database analysis. Cancer Letters, 2021, 522, 14-21.  | 3.2 | 10        |
| 11 | Predictive value of MGMT promoter methylation on the survival of TMZ treated <i>IDH</i> -mutant glioblastoma. Cancer Biology and Medicine, 2021, 18, 271-282.  | 1.4 | 31        |
| 12 | Transcriptional Characteristics of IDH-Wild Type Glioma Subgroups Highlight the Biological Processes Underlying Heterogeneity of IDH-Wild Type WHO Grade IV Gliomas. Frontiers in Cell and Developmental Biology, 2020, 8, 580464. | 1.8 | 8         |
| 13 | Identification of an ATP metabolismâ€related signature associated with prognosis and immune microenvironment in gliomas. Cancer Science, 2020, 111, 2325-2335.   | 1.7 | 27        |
| 14 | RGS16 promotes glioma progression and serves as a prognostic factor. CNS Neuroscience and Therapeutics, 2020, 26, 791-803.   | 1.9 | 24        |
| 15 | Combinations of four or more CpGs methylation present equivalent predictive value for MGMT expression and temozolomide therapeutic prognosis in gliomas. CNS Neuroscience and Therapeutics, 2019, 25, 314-322.                     | 1.9 | 42        |
| 16 | A Novel DNA Methylation-Based Signature Can Predict the Responses of MGMT Promoter Unmethylated Glioblastomas to Temozolomide. Frontiers in Genetics, 2019, 10, 910.   | 1.1 | 22        |
| 17 | Hypoxia induced LBH overexpression accelerates malignant progression in glioma. EBioMedicine, 2019, 49, 4-5.   | 2.7 | 1         |
| 18 | Systematically profiling the expression of eIF3 subunits in glioma reveals the expression of eIF3i has prognostic value in IDH-mutant lower grade glioma. Cancer Cell International, 2019, 19, 155.                                | 1.8 | 27        |

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|----|--|-----|-----------|
| 19 | Molecular and clinical characterization of TMEM71 expression at the transcriptional level in glioma. CNS Neuroscience and Therapeutics, 2019, 25, 965-975.   | 1.9 | 21        |
| 20 | Systematically characterize the clinical and biological significances of $1p19q$ genes in $1p/19q$ non-codeletion glioma. Carcinogenesis, $2019$ , $40$ , $1229$ - $1239$ .                                  | 1.3 | 60        |
| 21 | m6A RNA methylation regulators contribute to malignant progression and have clinical prognostic impact in gliomas. Aging, 2019, 11, 1204-1225.   | 1.4 | 209       |
| 22 | <p>Siglecs, Novel Immunotherapy Targets, Potentially Enhance The Effectiveness of Existing Immune Checkpoint Inhibitors in Glioma Immunotherapy</p> . OncoTargets and Therapy, 2019, Volume 12, 10263-10273. | 1.0 | 25        |
| 23 | ADAMTSL4, a Secreted Glycoprotein, Is a Novel Immune-Related Biomarker for Primary Glioblastoma<br>Multiforme. Disease Markers, 2019, 2019, 1-12.  | 0.6 | 66        |
| 24 | A novel analytical model of MGMT methylation pyrosequencing offers improved predictive performance in patients with gliomas. Modern Pathology, 2019, 32, 4-15.   | 2.9 | 41        |
| 25 | RNA processing genes characterize RNA splicing and further stratify lower-grade glioma. JCI Insight, 2019, 5, .  | 2.3 | 20        |
| 26 | A novel gene signature based on five glioblastoma stem-like cell relevant genes predicts the survival of primary glioblastoma. Journal of Cancer Research and Clinical Oncology, 2018, 144, 439-447.         | 1.2 | 36        |
| 27 | A comprehensive review of available omics data resources and molecular profiling for precision glioma studies (Review). Biomedical Reports, 2018, 10, 3-9.   | 0.9 | 7         |
| 28 | ADAR3 expression is an independent prognostic factor in lower-grade diffuse gliomas and positively correlated with the editing level of GRIA2Q607R. Cancer Cell International, 2018, 18, 196.                | 1.8 | 19        |
| 29 | Expression profile analysis of antisense long non-coding RNA identifies WDFY3-AS2 as a prognostic biomarker in diffuse glioma. Cancer Cell International, 2018, 18, 107.                                     | 1.8 | 33        |