

# Yu-Qing Liu

## List of Publications by Year in descending order

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Version: 2024-02-01

18  
papers

724  
citations

623574

14  
h-index

887953

17  
g-index

18  
all docs

18  
docs citations

18  
times ranked

832  
citing authors

#	ARTICLE	IF	CITATIONS
1	Recurrent PTPN11-MET fusion and a high occurrence rate of MET exon 14 skipping in brain metastases. <i>Cancer Science</i> , 2022, 113, 796-801.	1.7	7
2	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.	1.6	4
3	Molecular subtyping reveals immune alterations in IDH-wildtype lower-grade diffuse glioma. <i>Journal of Pathology</i> , 2020, 251, 272-283.	2.1	42
4	Classification of diffuse lower-grade glioma based on immunological profiling. <i>Molecular Oncology</i> , 2020, 14, 2081-2095.	2.1	48
5	The molecular characteristics of spinal cord gliomas with or without H3 K27M mutation. <i>Acta Neuropathologica Communications</i> , 2020, 8, 40.	2.4	51
6	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.	1.9	42
7	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.	1.1	22
8	Prognostic power of a lipid metabolism gene panel for diffuse gliomas. <i>Journal of Cellular and Molecular Medicine</i> , 2019, 23, 7741-7748.	1.6	59
9	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.	1.8	27
10	Systematically characterize the clinical and biological significances of 1p19q genes in 1p/19q non-codeletion glioma. <i>Carcinogenesis</i> , 2019, 40, 1229-1239.	1.3	60
11	Peripheral blood test provides a practical method for glioma evaluation and prognosis prediction. <i>CNS Neuroscience and Therapeutics</i> , 2019, 25, 876-883.	1.9	27
12	m6A RNA methylation regulators contribute to malignant progression and have clinical prognostic impact in gliomas. <i>Aging</i> , 2019, 11, 1204-1225.	1.4	209
13	Gene Expression Profiling Stratifies IDH-Wildtype Glioblastoma With Distinct Prognoses. <i>Frontiers in Oncology</i> , 2019, 9, 1433.	1.3	16
14	Amino acid metabolism-related gene expression-based risk signature can better predict overall survival for glioma. <i>Cancer Science</i> , 2019, 110, 321-333.	1.7	39
15	A novel analytical model of MGMT methylation pyrosequencing offers improved predictive performance in patients with gliomas. <i>Modern Pathology</i> , 2019, 32, 4-15.	2.9	41
16	RNA processing genes characterize RNA splicing and further stratify lower-grade glioma. <i>JCI Insight</i> , 2019, 5, .	2.3	20
17	Phosphohistone H3 (pHH3) is a prognostic and epithelial to mesenchymal transition marker in diffuse gliomas. <i>Oncotarget</i> , 2016, 7, 45005-45014.	0.8	10