Yu-Qing Liu

List of Publications by Year
in descending order

18	724	623574 14	887953 17
papers	citations	h-index	g-index

18	18	18	832
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	m6A RNA methylation regulators contribute to malignant progression and have clinical prognostic impact in gliomas. Aging, 2019, 11, 1204-1225.	1.4	209
2	Systematically characterize the clinical and biological significances of 1p19q genes in 1p/19q non-codeletion glioma. Carcinogenesis, 2019, 40, 1229-1239.	1.3	60
3	Prognostic power of a lipid metabolism gene panel for diffuse gliomas. Journal of Cellular and Molecular Medicine, 2019, 23, 7741-7748.	1.6	59
4	The molecular characteristics of spinal cord gliomas with or without H3 K27M mutation. Acta Neuropathologica Communications, 2020, 8, 40.	2.4	51
5	Classification of diffuse lowerâ€grade glioma based on immunological profiling. Molecular Oncology, 2020, 14, 2081-2095.	2.1	48
6	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
7	Molecular subtyping reveals immune alterations in <scp>IDH</scp> wildâ€type lowerâ€grade diffuse glioma. Journal of Pathology, 2020, 251, 272-283.	2.1	42
8	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
9	Amino acid metabolismâ€related gene expressionâ€based risk signature can better predict overall survival for glioma. Cancer Science, 2019, 110, 321-333.	1.7	39
10	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
11	Peripheral blood test provides a practical method for glioma evaluation and prognosis prediction. CNS Neuroscience and Therapeutics, 2019, 25, 876-883.	1.9	27
12	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
13	RNA processing genes characterize RNA splicing and further stratify lower-grade glioma. JCI Insight, 2019, 5, .	2.3	20
14	Gene Expression Profiling Stratifies IDH-Wildtype Glioblastoma With Distinct Prognoses. Frontiers in Oncology, 2019, 9, 1433.	1.3	16
15	Phosphohistone H3 (pHH3) is a prognostic and epithelial to mesenchymal transition marker in diffuse gliomas. Oncotarget, 2016, 7, 45005-45014.	0.8	10
16	Recurrent PTPRZ1â€MET fusion and a high occurrence rate of MET exon 14 skipping in brain metastases. Cancer Science, 2022, 113, 796-801.	1.7	7
17	Characterization and prognostic significance of alternative splicing events in lowerâ€grade diffuse gliomas. Journal of Cellular and Molecular Medicine, 2020, 24, 13171-13180.	1.6	4
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