

Rose K Lai

List of Publications by Year in descending order

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Version: 2024-02-01

17
papers

907
citations

687363

13
h-index

888059

17
g-index

18
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18
docs citations

18
times ranked

2165
citing authors

#	ARTICLE	IF	CITATIONS
1	A functional variant on 20q13.33 related to glioma risk alters enhancer activity and modulates expression of multiple genes. <i>Human Mutation</i> , 2021, 42, 77-88.	2.5	12
2	Phase I trial of intranasal NEO100, highly purified perillyl alcohol, in adult patients with recurrent glioblastoma. <i>Neuro-Oncology Advances</i> , 2021, 3, vdab005.	0.7	8
3	A Functional Variant on 9p21.3 Related to Glioma Risk Affects Enhancer Activity and Modulates Expression of <i>CDKN2B</i> and <i>AS1</i> . <i>Human Mutation</i> , 2021, 42, 1208-1214.	2.5	8
4	Meta-Analyses of Splicing and Expression Quantitative Trait Loci Identified Susceptibility Genes of Glioma. <i>Frontiers in Genetics</i> , 2021, 12, 609657.	2.3	1
5	Lack of association between modifiable exposures and glioma risk: A Mendelian randomisation analysis. <i>Neuro-Oncology</i> , 2020, 22, 207-215.	1.2	19
6	Glioma risk associated with extent of estimated European genetic ancestry in African Americans and Hispanics. <i>International Journal of Cancer</i> , 2020, 146, 739-748.	5.1	23
7	Aspirin, NSAIDs, and Glioma Risk: Original Data from the Glioma International Case-Control Study and a Meta-analysis. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2019, 28, 555-562.	2.5	15
8	Sex-specific gene and pathway modeling of inherited glioma risk. <i>Neuro-Oncology</i> , 2019, 21, 71-82.	1.2	52
9	Transcriptome-Wide Association Study Identifies New Candidate Susceptibility Genes for Glioma. <i>Cancer Research</i> , 2019, 79, 2065-2071.	0.9	26
10	Mendelian randomisation study of the relationship between vitamin D and risk of glioma. <i>Scientific Reports</i> , 2018, 8, 2339.	3.3	23
11	Impact of atopy on risk of glioma: a Mendelian randomisation study. <i>BMC Medicine</i> , 2018, 16, 42.	5.5	38
12	Influence of obesity-related risk factors in the aetiology of glioma. <i>British Journal of Cancer</i> , 2018, 118, 1020-1027.	6.4	32
13	Age-specific genome-wide association study in glioblastoma identifies increased proportion of lower grade glioma-like features associated with younger age. <i>International Journal of Cancer</i> , 2018, 143, 2359-2366.	5.1	21
14	Sex-specific glioma genome-wide association study identifies new risk locus at 3p21.31 in females, and finds sex-differences in risk at 8q24.21. <i>Scientific Reports</i> , 2018, 8, 7352.	3.3	56
15	Genome-wide association study of glioma subtypes identifies specific differences in genetic susceptibility to glioblastoma and non-glioblastoma tumors. <i>Nature Genetics</i> , 2017, 49, 789-794.	21.4	259
16	Circular RNA profile in gliomas revealed by identification tool UROBORUS. <i>Nucleic Acids Research</i> , 2016, 44, e87-e87.	14.5	269
17	Genome-Wide Methylation Analyses in Glioblastoma Multiforme. <i>PLoS ONE</i> , 2014, 9, e89376.	2.5	45