

Ying-Chih Wang

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9088706/publications.pdf>

Version: 2024-02-01

18
papers

2,701
citations

1040056

9
h-index

1125743

13
g-index

19
all docs

19
docs citations

19
times ranked

7073
citing authors

#	ARTICLE	IF	CITATIONS
1	Abstract P046: NKG2A and HLA-E define a novel alternative immune checkpoint axis in bladder cancer. , 2022, , .		0
2	Molecular evidence of SARS-CoV-2 in New York before the first pandemic wave. Nature Communications, 2021, 12, 3463.	12.8	18
3	Downregulation of exhausted cytotoxic T cells in gene expression networks of multisystem inflammatory syndrome in children. Nature Communications, 2021, 12, 4854.	12.8	42
4	621â€¦NKG2A and HLA-E define a novel mechanism of resistance to immunotherapy with M. bovis BCG in non-muscle-invasive bladder cancer patients. , 2021, 9, A651-A651.		0
5	314â€¦NKG2A and HLA-E define a novel alternative immune checkpoint axis in bladder cancer. , 2021, 9, A338-A338.		1
6	Negative feedback control of neuronal activity by microglia. Nature, 2020, 586, 417-423.	27.8	520
7	Multiscale causal networks identify VGF as a key regulator of Alzheimerâ€™s disease. Nature Communications, 2020, 11, 3942.	12.8	94
8	Introductions and early spread of SARS-CoV-2 in the New York City area. Science, 2020, 369, 297-301.	12.6	356
9	Recurrent IDH mutations in high-grade meningioma. Neuro-Oncology, 2020, 22, 1044-1045.	1.2	10
10	Truncated stathmin-2 is a marker of TDP-43 pathology in frontotemporal dementia. Journal of Clinical Investigation, 2020, 130, 6080-6092.	8.2	117
11	EPCO-20. RELATING GLIOBLASTOMA HETEROGENEITY TO HUMAN FETAL GLIAL DEVELOPMENT THROUGH HIGH RESOLUTION SINGLE NUCLEI TRANSCRIPTOMICS. Neuro-Oncology, 2020, 22, ii73-ii73.	1.2	0
12	O1.3. DIFFERENTIAL HISTONE MODIFICATIONS IN 250 SCHIZOPHRENIA CASES AND 330 CONTROLS. Schizophrenia Bulletin, 2019, 45, S159-S160.	4.3	0
13	CommonMind Consortium provides transcriptomic and epigenomic data for Schizophrenia and Bipolar Disorder. Scientific Data, 2019, 6, 180.	5.3	149
14	High-coverage, long-read sequencing of Han Chinese trio reference samples. Scientific Data, 2019, 6, 91.	5.3	13
15	Integrative transcriptome analyses of the aging brain implicate altered splicing in Alzheimerâ€™s disease susceptibility. Nature Genetics, 2018, 50, 1584-1592.	21.4	307
16	Cell-specific histone modification maps in the human frontal lobe link schizophrenia risk to the neuronal epigenome. Nature Neuroscience, 2018, 21, 1126-1136.	14.8	112
17	Gene expression elucidates functional impact of polygenic risk for schizophrenia. Nature Neuroscience, 2016, 19, 1442-1453.	14.8	952
18	The somatic mutational landscape in soft tissue sarcoma: Early results from TCGA data.. Journal of Clinical Oncology, 2015, 33, 10508-10508.	1.6	8