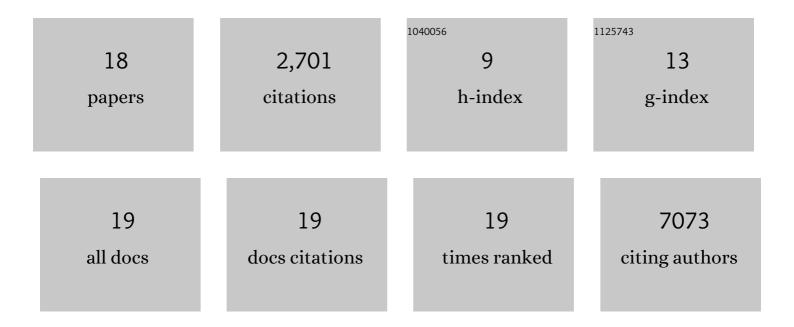
## Ying-Chih Wang

List of Publications by Year in descending order

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YING-CHIH WANG

#	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 Alzheimer'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 Alzheimer'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