Hung-Hsin Chen

List of Publications by Year
in descending order

23	410	933264 10	794469 19
papers	citations	h-index	g-index

24	24	24	991
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	The metabolome profiling and pathway analysis in metabolic healthy and abnormal obesity. International Journal of Obesity, 2015, 39, 1241-1248.	1.6	107
2	Hepatitis C virus infection: a risk factor for <scp>P</scp>arkinson's disease. Journal of Viral Hepatitis, 2015, 22, 784-791.	1.0	75
3	The hypoxia-responsive lncRNANDRG-OT1promotes NDRG1 degradation via ubiquitin-mediated proteolysis in breast cancer cells. Oncotarget, 2018, 9, 10470-10482.	0.8	33
4	Body mass index and breast cancer: analysis of a nation-wide population-based prospective cohort study on 1 393 985 Taiwanese women. International Journal of Obesity, 2016, 40, 524-530.	1.6	32
5	A dynamic model for the outbreaks of hand, foot, and mouth disease in Taiwan. Epidemiology and Infection, 2016, 144, 1500-1511.	1.0	26
6	Host genetic effects in pneumonia. American Journal of Human Genetics, 2021, 108, 194-201.	2.6	17
7	Identifying developmental stuttering and associated comorbidities in electronic health records and creating a phenome risk classifier. Journal of Fluency Disorders, 2021, 68, 105847.	0.7	17
8	Genetically regulated expression in late-onset Alzheimerâ€™s disease implicates risk genes within known and novel loci. Translational Psychiatry, 2021, 11, 618.	2.4	17
9	Irritable bowel syndrome and the incidence of colorectal neoplasia: a prospective cohort study with community-based screened population in Taiwan. British Journal of Cancer, 2015, 112, 171-176.	2.9	16
10	Functionally oriented analysis of cardiometabolic traits in a trans-ethnic sample. Human Molecular Genetics, 2019, 28, 1212-1224.	1.4	12
11	A pulmonary tuberculosis outbreak in a long-term care facility. Epidemiology and Infection, 2016, 144, 1455-1462.	1.0	11
12	Phenome risk classification enables phenotypic imputation and gene discovery in developmental stuttering. American Journal of Human Genetics, 2021, 108, 2271-2283.	2.6	11
13	Estimating the benefits of adjuvant systemic therapy for women with early breast cancer. British Journal of Surgery, 2002, 88, 1513-1518.	0.1	10
14	Effects of risk factors on periodontal disease defined by calibrated community periodontal index and loss of attachment scores. Oral Diseases, 2017, 23, 949-955.	1.5	7
15	GWAS and Beyond: Using Omics Approaches to Interpret SNP Associations. Current Genetic Medicine Reports, 2019, 7, 30-40.	1.9	4
16	Population-based genetic effects for developmental stuttering. Human Genetics and Genomics Advances, 2022, 3, 100073.	1.0	4
17	Novel diabetes gene discovery through comprehensive characterization and integrative analysis of longitudinal gene expression changes. Human Molecular Genetics, 2022, 31, 3191-3205.	1.4	4
18	Risk factors for incidence and case-fatality rates of healthcare-associated infections: a 20-year follow-up of a hospital-based cohort. Epidemiology and Infection, 2016, 144, 198-206.	1.0	3

#	Article	IF	Citations
19	Ancestral diversity improves discovery and fine-mapping of genetic loci for anthropometric traitsâ€"The Hispanic/Latino Anthropometry Consortium. Human Genetics and Genomics Advances, 2022, 3, 100099.	1.0	3
20	Bone mineral density as a dose-response predictor for osteoporosis: a propensity score analysis of longitudinal incident study (KCIS no. 39). QJM - Monthly Journal of the Association of Physicians, 2019, 112, 327-333.	0.2	1
21	Optimizing Genetic Analyses of Serum Lipids in Longitudinal Data. Circulation Research, 2020, 127, 1337-1339.	2.0	0
22	Tissueâ€specific genetically regulated expression in lateâ€onset Alzheimerâ€™s disease implicates risk genes within known and 30 novel loci. Alzheimer's and Dementia, 2020, 16, e039475.	0.4	0
23	Reduction of Colorectal Cancer Mortality and Advanced Stage Cancer Incidence After 10 Years of Fecal Immunochemical Test Screening. Journal of Global Oncology, 2018, 4, 206s-206s.	0.5	0