Yen-Feng Chiu

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4000957/publications.pdf

Version: 2024-02-01

46 papers

1,241 citations

16 h-index 32 g-index

48 all docs

48 docs citations

48 times ranked

2547 citing authors

#	Article	IF	CITATIONS
1	Assessment of incident frailty hazard associated with depressive symptoms in a Taiwanese longitudinal study. International Psychogeriatrics, 2022, 34, 61-70.	1.0	9
2	A phase II randomised trial of induction chemotherapy followed by concurrent chemoradiotherapy in locally advanced pancreatic cancer: the Taiwan Cooperative Oncology Group T2212 study. British Journal of Cancer, 2022, 126, 1018-1026.	6.4	11
3	Multi-omics analysis identifies CpGs near G6PC2 mediating the effects of genetic variants on fasting glucose. Diabetologia, 2021, 64, 1613-1625.	6.3	9
4	The trans-ancestral genomic architecture of glycemic traits. Nature Genetics, 2021, 53, 840-860.	21.4	341
5	Longitudinal changes in physical and mental health of older adults with chronic hepatitis B infection: Trajectories and predictors. Preventive Medicine Reports, 2021, 23, 101432.	1.8	1
6	The paradox of aging and health-related quality of life in Asian Chinese: results from the Healthy Aging Longitudinal Study in Taiwan. BMC Geriatrics, 2020, 20, 91.	2.7	12
7	Associations of autozygosity with a broad range of human phenotypes. Nature Communications, 2019, 10, 4957.	12.8	84
8	Genomeâ€wide scan for circulating vascular adhesion protein†levels: <i><scp>MACROD</scp>2</i> as a potential transcriptional regulator of adipogenesis. Journal of Diabetes Investigation, 2018, 9, 1067-1074.	2.4	13
9	Sedative–hypnotic drug use among community-dwelling elderly in Taiwan. International Psychogeriatrics, 2018, 30, 957-965.	1.0	7
10	Assessment of fenofibrate-methylation interactions on triglycerides using longitudinal family data. BMC Proceedings, 2018, 12, 48.	1.6	3
11	General retrospective megaâ€analysis framework for rare variant association tests. Genetic Epidemiology, 2018, 42, 621-635.	1.3	O
12	The Role of Consolidation Chemoradiotherapy in Locally Advanced Pancreatic Cancer Receiving Chemotherapy: An Updated Systematic Review and Meta-Analysis. Cancer Research and Treatment, 2018, 50, 562-574.	3.0	16
13	Independent association between subjective cognitive decline and frailty in the elderly. PLoS ONE, 2018, 13, e0201351.	2.5	33
14	Moderating effect of depression on the association between pain and activities of daily living in older adults. Psychogeriatrics, 2018, 18, 379-387.	1.2	6
15	<i>IGF1</i> Gene Is Associated With Triglyceride Levels In Subjects With Family History Of Hypertension From The SAPPHIRe And TWB Projects. International Journal of Medical Sciences, 2018, 15, 1035-1042.	2.5	3
16	Prevalence of and risk factors for minor and major depression among community-dwelling older adults in Taiwan. International Psychogeriatrics, 2017, 29, 1113-1121.	1.0	31
17	Regionâ€based association tests for sequencing data on survival traits. Genetic Epidemiology, 2017, 41, 511-522.	1.3	6
18	Cohort Profile: The Healthy Aging Longitudinal Study in Taiwan (HALST). International Journal of Epidemiology, 2017, 46, 1106-1106j.	1.9	26

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19	Genome-wide copy number variation analysis identified deletions in SFMBT1 associated with fasting plasma glucose in a Han Chinese population. BMC Genomics, 2017, 18, 591.	2.8	8
20	Generalization of Rare Variant Association Tests for Longitudinal Family Studies. Genetic Epidemiology, 2016, 40, 101-112.	1.3	6
21	Vegetarian diet reduces the risk of hypertension independent of abdominal obesity and inflammation. Journal of Hypertension, 2016, 34, 2164-2171.	0.5	47
22	Interleukin-6 gene polymorphisms correlate with the progression of nephropathy in Chinese patients with type 2 diabetes: A prospective cohort study. Diabetes Research and Clinical Practice, 2016, 120, 15-23.	2.8	15
23	Multipoint association mapping for longitudinal family data: an application to hypertension phenotypes. BMC Proceedings, 2016, 10, 315-320.	1.6	3
24	Longitudinal analytical approaches to genetic data. BMC Genetics, 2016, 17, 4.	2.7	19
25	Genetics of Coronary Artery Disease in Taiwan: A Cardiometabochip Study by the Taichi Consortium. PLoS ONE, 2016, 11, e0138014.	2.5	33
26	Cross-sectional and longitudinal comparisons of metabolic profiles between vegetarian and non-vegetarian subjects: a matched cohort study. British Journal of Nutrition, 2015, 114, 1313-1320.	2.3	114
27	Simultaneous estimation of the locations and effects of multiple disease loci in case-control studies. Biostatistics, 2015, 16, 222-239.	1.5	0
28	A meta-analysis of genome-wide association studies for adiponectin levels in East Asians identifies a novel locus near WDR11-FGFR2. Human Molecular Genetics, 2014, 23, 1108-1119.	2.9	68
29	Adiponectin gene (ADIPOQ) polymorphisms correlate with the progression of nephropathy in Taiwanese male patients with type 2 diabetes. Diabetes Research and Clinical Practice, 2014, 105, 261-270.	2.8	17
30	Identification of rare variants for hypertension with incorporation of linkage information. BMC Proceedings, 2014, 8, S109.	1.6	5
31	Taiwanese Vegetarians and Omnivores: Dietary Composition, Prevalence of Diabetes and IFG. PLoS ONE, 2014, 9, e88547.	2.5	67
32	Analysis of Family―and Populationâ€Based Samples Using Multiple Linkage Disequilibrium Mapping. Annals of Human Genetics, 2013, 77, 251-267.	0.8	2
33	A Novel Support Vector Machine-Based Approach for Rare Variant Detection. PLoS ONE, 2013, 8, e71114.	2.5	4
34	SVMâ€Based Generalized Multifactor Dimensionality Reduction Approaches for Detecting Geneâ€Gene Interactions in Family Studies. Genetic Epidemiology, 2012, 36, 88-98.	1.3	22
35	Incorporation of covariates in simultaneous localization of two linked loci using affected relative pairs. BMC Genetics, 2010, 11, 67.	2.7	2
36	SLC2A10 genetic polymorphism predicts development of peripheral arterial disease in patients with type 2 diabetes. SLC2A10and PAD in type 2 diabetes. BMC Medical Genetics, 2010, 11, 126.	2.1	15

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37	Incorporating Covariates into Multipoint Association Mapping in the Case-Parent Design. Human Heredity, 2010, 69, 229-241.	0.8	4
38	Assessment of gene-covariate interactions by incorporating covariates into association mapping. BMC Proceedings, 2009, 3, S85.	1.6	3
39	Incorporation of covariates into multipoint linkage disequilibrium mapping in caseâ€control studies. Genetic Epidemiology, 2008, 32, 143-151.	1.3	4
40	Incorporating quantitative variables into linkage analysis using affected sib pairs. BMC Proceedings, 2007, 1, S98.	1.6	2
41	Multipoint linkage mapping using sibpairs: Non-parametric estimation of trait effects with quantitative covariates. Genetic Epidemiology, 2005, 28, 58-69.	1.3	8
42	Multipoint linkage disequilibrium mapping using case-control designs. Genetic Epidemiology, 2005, 29, 365-376.	1.3	8
43	A comparison in association and linkage genome-wide scans for alcoholism susceptibility genes using single-nucleotide polymorphisms. BMC Genetics, 2005, 6, S89.	2.7	5
44	An Autosomal Genome-wide Scan for Loci Linked to Pre-Diabetic Phenotypes in Nondiabetic Chinese Subjects From the Stanford Asia-Pacific Program of Hypertension and Insulin Resistance Family Study. Diabetes, 2005, 54, 1200-1206.	0.6	29
45	Multipoint affected sibpair linkage methods for localizing susceptibility genes of complex diseases. Genetic Epidemiology, 2003, 24, 107-117.	1.3	33
46	A Robust Identity-by-Descent Procedure Using Affected Sib Pairs: Multipoint Mapping for Complex Diseases. Human Heredity. 2001. 51. 64-78.	0.8	67