Jingyun Yang

List of Publications by Year
in descending order

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		201674	133252
62	4,690	27	59
papers	citations	h-index	g-index

65	65	65	9791
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Novel DNA methylation loci and genes showing pleiotropic association with Alzheimerâ€™s dementia: a network Mendelian randomization analysis. Epigenetics, 2022, 17, 746-758.	2.7	8
2	Mendelian Randomization Analysis Identified Potential Genes Pleiotropically Associated with Polycystic Ovary Syndrome. Reproductive Sciences, 2022, 29, 1028-1037.	2.5	6
3	Neuropathologic Correlates of Human Cortical Proteins in Alzheimer Disease and Related Dementias. Neurology, 2022, 98, .	1.1	9
4	0277 Deep learning revealed associations between altered temporal correlations in motor activity and Parkinsonâ€™s risk. Sleep, 2022, 45, A124-A125.	1.1	0
5	Mendelian randomization integrating GWAS and mQTL data identified novel pleiotropic DNA methylation loci for neuropathology of Alzheimerâ€™s disease. Neurobiology of Aging, 2021, 97, 18-27.	3.1	14
6	Mendelian randomization analysis identified genes pleiotropically associated with the risk and prognosis of COVID-19. Journal of Infection, 2021, 82, 126-132.	3.3	37
7	Association of sleep disturbance and freezing of gait in Parkinson disease: prevention/delay implications. Journal of Clinical Sleep Medicine, 2021, 17, 779-789.	2.6	10
8	Monitoring recessions: A Bayesian sequential quickest detection method. International Journal of Forecasting, 2021, 37, 500-510.	6.5	3
9	Genome-wide meta-analysis of muscle weakness identifies 15 susceptibility loci in older men and women. Nature Communications, 2021, 12, 654.	12.8	75
10	Systemic brain derived neurotrophic factor but not intestinal barrier integrity is associated with cognitive decline and incident Alzheimerâ€™s disease. PLoS ONE, 2021, 16, e0240342.	2.5	6
11	Deep learningâ€based detection and stage grading for optimising diagnosis of diabetic retinopathy. Diabetes/Metabolism Research and Reviews, 2021, 37, e3445.	4.0	16
12	Mendelian randomization integrating GWAS and eQTL data revealed genes pleiotropically associated with major depressive disorder. Translational Psychiatry, 2021, 11, 225.	4.8	19
13	Cognitive and brain cytokine profile of non-demented individuals with cerebral amyloid-beta deposition. Journal of Neuroinflammation, 2021, 18, 147.	7.2	11
14	Mendelian randomization analysis identified genes potentially pleiotropically associated with periodontitis. Saudi Journal of Biological Sciences, 2021, 28, 4089-4095.	3.8	4
15	Mendelian randomization analysis identified genes pleiotropically associated with central corneal thickness. BMC Genomics, 2021, 22, 517.	2.8	9
16	Bootstrap approach for meta-synthesis of MRI findings from multiple scanners. Journal of Neuroscience Methods, 2021, 360, 109229.	2.5	1
17	Latent Cognitive Class at Enrollment Predicts Future Cognitive Trajectories of Decline in a Community Sample of Older Adults. Journal of Alzheimer's Disease, 2021, 83, 641-652.	2.6	1
18	Human Brain and Blood N-Glycome Profiling in Alzheimerâ€™s Disease and Alzheimerâ€™s Disease-Related Dementias. Frontiers in Aging Neuroscience, 2021, 13, 765259.	3.4	8

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19	The power of genetic diversity in genome-wide association studies of lipids. Nature, 2021, 600, 675-679.	27.8	353
20	A meta-analysis of genome-wide association studies identifies new genetic loci associated with all-cause and vascular dementia.. Alzheimer's and Dementia, 2021, 17 Suppl 3, e056081.	0.8	0
21	Brain and blood metabolome for Alzheimer's dementia: findings from a targeted metabolomics analysis. Neurobiology of Aging, 2020, 86, 123-133.	3.1	83
22	On the predictive performance of two Bayesian joint models: a simulation study. Communications in Statistics Part B: Simulation and Computation, 2020, , 1-10.	1.2	1
23	The association between genetic variants in lactotransferrin and dental caries: a meta- and gene-based analysis. BMC Medical Genetics, 2020, 21, 114.	2.1	4
24	Genome-wide interaction analysis of pathological hallmarks in Alzheimer's disease. Neurobiology of Aging, 2020, 93, 61-68.	3.1	63
25	Associations of autozygosity with a broad range of human phenotypes. Nature Communications, 2019, 10, 4957.	12.8	84
26	On the performance of MixTVEM: a simulation study. Communications in Statistics Part B: Simulation and Computation, 2019, 48, 2830-2844.	1.2	2
27	Interaction between the progression of Alzheimer's disease and fractal degradation. Neurobiology of Aging, 2019, 83, 21-30.	3.1	22
28	Postmortem neurodegenerative markers and trajectories of decline in cognitive systems. Neurology, 2019, 92, e831-e840.	1.1	34
29	Genetic architecture of subcortical brain structures in 38,851 individuals. Nature Genetics, 2019, 51, 1624-1636.	21.4	192
30	DNA methylation variability in Alzheimer's disease. Neurobiology of Aging, 2019, 76, 35-44.	3.1	25
31	Genome-wide association study of 23,500 individuals identifies 7 loci associated with brain ventricular volume. Nature Communications, 2018, 9, 3945.	12.8	31
32	Study of 300,486 individuals identifies 148 independent genetic loci influencing general cognitive function. Nature Communications, 2018, 9, 2098.	12.8	484
33	Neurodegenerative disease and cognitive retest learning. Neurobiology of Aging, 2018, 66, 122-130.	3.1	10
34	Novel genetic loci associated with hippocampal volume. Nature Communications, 2017, 8, 13624.	12.8	250
35	A genomeâ€wide profiling of brain DNA hydroxymethylation in Alzheimer's disease. Alzheimer's and Dementia, 2017, 13, 674-688.	0.8	83
36	Neuroprotective effects respond to cerebral ischemia without susceptibility to HBâ€tumorigenesis in VHL heterozygous knockout mice. Molecular Carcinogenesis, 2017, 56, 2342-2351.	2.7	2

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37	An Analysis of Two Genome-wide Association Meta-analyses Identifies a New Locus for Broad Depression Phenotype. Biological Psychiatry, 2017, 82, 322-329.	1.3	84
38	[P4â€“035]: AMYLOID Î²â€DRIVEN DNA DEMETHYLATION AS A TARGET FOR ALZHEIMER's DISEASE. Alzheimer's and Dementia, 2017, 13, P1269.	0.8	0
39	Association Between Brain Gene Expression, DNA Methylation, and Alteration of Ex Vivo Magnetic Resonance Imaging Transverse Relaxation in Late-Life Cognitive Decline. JAMA Neurology, 2017, 74, 1473.	9.0	21
40	Association of Bone Metabolic Markers With Diabetic Retinopathy and Diabetic Macular Edema in Elderly Chinese Individuals With Type 2 Diabetes Mellitus. American Journal of the Medical Sciences, 2017, 354, 355-361.	1.1	8
41	Genome-Wide Association Analysis of the Sense of Smell in U.S. Older Adults: Identification of Novel Risk Loci in African-Americans and European-Americans. Molecular Neurobiology, 2017, 54, 8021-8032.	4.0	17
42	Identification of genes associated with dissociation of cognitive performance and neuropathological burden: Multistep analysis of genetic, epigenetic, and transcriptional data. PLoS Medicine, 2017, 14, e1002287.	8.4	88
43	Varied effects of age-related neuropathologies on the trajectory of late life cognitive decline. Brain, 2017, 140, aww341.	7.6	81
44	APOE Îµ4-TOMM40 â€˜523 haplotypes and the risk of Alzheimerâ€™s disease in older Caucasian and African Americans. PLoS ONE, 2017, 12, e0180356.	2.5	39
45	Personality Polygenes, Positive Affect, and Life Satisfaction. Twin Research and Human Genetics, 2016, 19, 407-417.	0.6	16
46	O2â€02â€01: Dna Demethylation and Remethylation in Alzheimerâ€™s Pathology. Alzheimer's and Dementia, 2016, 12, P223.	0.8	0
47	Methylation profiles in peripheral blood CD4+ lymphocytes versus brain: The relation to Alzheimer's disease pathology. Alzheimer's and Dementia, 2016, 12, 942-951.	0.8	44
48	Genome-wide association study identifies 74 loci associated with educational attainment. Nature, 2016, 533, 539-542.	27.8	1,204
49	Novel genetic loci underlying human intracranial volume identified through genome-wide association. Nature Neuroscience, 2016, 19, 1569-1582.	14.8	213
50	Rescue of Early bace-1 and Global DNA Demethylation by S-Adenosylmethionine Reduces Amyloid Pathology and Improves Cognition in an Alzheimerâ€™s Model. Scientific Reports, 2016, 6, 34051.	3.3	49
51	Genome-wide association analysis identifies genetic loci associated with resistance to multiple antimalarials in Plasmodium falciparum from China-Myanmar border. Scientific Reports, 2016, 6, 33891.	3.3	100
52	Genetic variants linked to education predict longevity. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 13366-13371.	7.1	110
53	CD41 and CD45 expression marks the angioformative initiation of neovascularisation in human haemangioblastoma. Tumor Biology, 2016, 37, 3765-3774.	1.8	11
54	Early life instruction in foreign language and music and incidence of mild cognitive impairment.. Neuropsychology, 2015, 29, 292-302.	1.3	75

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55	Aberrantly expressed mRNAs and long non-coding RNAs in patients with invasive ductal breast carcinoma: A pilot study. Molecular Medicine Reports, 2015, 11, 2185-2190.	2.4	7
56	Measuring Disagreement in Qualitative Expectations. Journal of Forecasting, 2015, 34, 405-426.	2.8	34
57	Physical activity, motor function, and white matter hyperintensity burden in healthy older adults. Neurology, 2015, 84, 1294-1300.	1.1	67
58	The TMEM106B locus and TDP-43 pathology in older persons without FTLD. Neurology, 2015, 84, 927-934.	1.1	71
59	Association of Brain DNA Methylation inSORL1,ABCA7,HLA-DRB5,SLC24A4, andBIN1With Pathological Diagnosis of Alzheimer Disease. JAMA Neurology, 2015, 72, 15.	9.0	239
60	Association of DNA methylation in the brain with age in older persons is confounded by common neuropathologies. International Journal of Biochemistry and Cell Biology, 2015, 67, 58-64.	2.8	34
61	Epigenomics of Alzheimer's disease. Translational Research, 2015, 165, 200-220.	5.0	97
62	Joint Association of Nicotinic Acetylcholine Receptor Variants with Abdominal Obesity in American Indians: The Strong Heart Family Study. PLoS ONE, 2014, 9, e102220.	2.5	10