## Yunzhang Wang

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7738777/publications.pdf

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623734 552781 1,116 26 14 26 citations g-index h-index papers 31 31 31 1657 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Longitudinal trajectories, correlations and mortality associations of nine biological ages across 20-years follow-up. ELife, 2020, 9, .	6.0	177
2	Within-sibship genome-wide association analyses decrease bias in estimates of direct genetic effects. Nature Genetics, 2022, 54, 581-592.	21.4	142
3	Age, Frailty, and Comorbidity as Prognostic Factors for Short-Term Outcomes in Patients With Coronavirus Disease 2019 in Geriatric Care. Journal of the American Medical Directors Association, 2020, 21, 1555-1559.e2.	2.5	141
4	A computational solution for bolstering reliability of epigenetic clocks: implications for clinical trials and longitudinal tracking. Nature Aging, 2022, 2, 644-661.	11.6	95
5	Genome-wide association studies identify 137 genetic loci for DNA methylation biomarkers of aging. Genome Biology, 2021, 22, 194.	8.8	90
6	A genomeâ€wide association study of the frailty index highlights brain pathways in ageing. Aging Cell, 2021, 20, e13459.	6.7	74
7	Epigenetic influences on aging: a longitudinal genome-wide methylation study in old Swedish twins. Epigenetics, 2018, 13, 975-987.	2.7	65
8	Deciphering the genetic and epidemiological landscape of mitochondrial DNA abundance. Human Genetics, 2021, 140, 849-861.	3.8	47
9	Human aging DNA methylation signatures are conserved but accelerated in cultured fibroblasts. Epigenetics, 2019, 14, 961-976.	2.7	36
10	Frailty and comorbidity in predicting community <scp>COVID</scp> â€19 mortality in the <scp>U.K.</scp> Biobank: The effect of sampling. Journal of the American Geriatrics Society, 2021, 69, 1128-1139.	2.6	32
11	DNA Methylation and All-Cause Mortality in Middle-Aged and Elderly Danish Twins. Genes, 2018, 9, 78.	2.4	27
12	Apolipoprotein E DNA methylation and late-life disease. International Journal of Epidemiology, 2018, 47, 899-907.	1.9	22
13	Comprehensive longitudinal study of epigenetic mutations in aging. Clinical Epigenetics, 2019, 11, 187.	4.1	21
14	DNA methylation signatures of aggression and closely related constructs: A meta-analysis of epigenome-wide studies across the lifespan. Molecular Psychiatry, 2021, 26, 2148-2162.	7.9	21
15	Genetically-predicted life-long lowering of low-density lipoprotein cholesterol is associated with decreased frailty: A Mendelian randomization study in UK biobank. EBioMedicine, 2019, 45, 487-494.	6.1	19
16	DNA methylation outlier burden, health, and ageing in Generation Scotland and the Lothian Birth Cohorts of 1921 and 1936. Clinical Epigenetics, 2020, 12, 49.	4.1	17
17	Frailty trajectories in three longitudinal studies of aging: Is the level or the rate of change more predictive of mortality?. Age and Ageing, 2021, 50, 2174-2182.	1.6	16
18	Implementing a method for studying longitudinal DNA methylation variability in association with age. Epigenetics, 2018, 13, 866-874.	2.7	13

#	Article	IF	CITATIONS
19	Clinical biomarkers and associations with healthspan and lifespan: Evidence from observational and genetic data. EBioMedicine, 2021, 66, 103318.	6.1	12
20	Frailty and the risk of dementia: is the association explained by shared environmental and genetic factors?. BMC Medicine, 2021, 19, 248.	5.5	11
21	Profiles of histidine-rich glycoprotein associate with age and risk of all-cause mortality. Life Science Alliance, 2020, 3, e202000817.	2.8	9
22	Fatty Acids and Frailty: A Mendelian Randomization Study. Nutrients, 2021, 13, 3539.	4.1	8
23	Replicating associations between DNA methylation and body mass index in a longitudinal sample of older twins. International Journal of Obesity, 2020, 44, 1397-1405.	3.4	6
24	The epigenetic etiology of cardiovascular disease in a longitudinal Swedish twin study. Clinical Epigenetics, 2021, 13, 129.	4.1	6
25	Protein Nutritional Status and Frailty: A Mendelian Randomization Study. Journal of Nutrition, 2022, 152, 269-275.	2.9	4
26	Epigenome-wide association study of level and change in cognitive abilities from midlife through late life. Clinical Epigenetics, 2021, 13, 85.	4.1	0