Xu Gao

List of Publications by Year in descending order

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289141 361296 47 1,886 20 40 citations h-index g-index papers 52 52 52 2983 all docs docs citations times ranked citing authors

#	Article	IF	CITATIONS
1	DunedinPACE, a DNA methylation biomarker of the pace of aging. ELife, 2022, 11, .	2.8	214
2	Epigenetics of early-life adversity in youth: cross-sectional and longitudinal associations. Clinical Epigenetics, 2022, 14, 48.	1.8	8
3	Associations of Carbonaceous Compounds and Water-Soluble Inorganic Ions in Ambient PM _{2.5} with Renal Function in Older Individuals: The China BAPE Study. Environmental Science & Earny; Technology, 2022, 56, 433-439.	4.6	4
4	Relationship between DNA Methylation Profiles and Active Tuberculosis Development from Latent Infection: a Pilot Study in Nested Case-Control Design. Microbiology Spectrum, 2022, 10, e0058622.	1.2	3
5	Role of sleep quality in the acceleration of biological aging and its potential for preventive interaction on air pollution insults: Findings from the UK Biobank cohort. Aging Cell, 2022, 21, e13610.	3.0	25
6	Association of DNA methylation in circulating CD4+T cells with short-term PM2.5 pollution waves: A quasi-experimental study of healthy young adults. Ecotoxicology and Environmental Safety, 2022, 239, 113634.	2.9	3
7	The Effect of Built Environment on Physical Health and Mental Health of Adults: A Nationwide Cross-Sectional Study in China. International Journal of Environmental Research and Public Health, 2022, 19, 6492.	1.2	5
8	Epigenetic age stratifies the risk of blood pressure elevation related to short-term PM2.5 exposure in older adults. Environmental Research, 2022, 212, 113507.	3.7	5
9	Mortality and morbidity risk prediction for older former smokers based on a score of smoking history: evidence from UK Biobank and ESTHER cohorts. Age and Ageing, 2022, 51, .	0.7	2
10	Associations of Plasma Folate and Vitamin B6 With Blood DNA Methylation Age: An Analysis of One-Carbon Metabolites in the VA Normative Aging Study. Journals of Gerontology - Series A Biological Sciences and Medical Sciences, 2021, 76, 760-769.	1.7	11
11	DNA methylation-based biomarkers of age acceleration and all-cause death, myocardial infarction, stroke, and cancer in two cohorts: The NAS, and KORA F4. EBioMedicine, 2021, 63, 103151.	2.7	42
12	Short-term air pollution, cognitive performance and nonsteroidal anti-inflammatory drug use in the Veterans Affairs Normative Aging Study. Nature Aging, 2021, 1, 430-437.	5. 3	33
13	DNAm-based signatures of accelerated aging and mortality in blood are associated with low renal function. Clinical Epigenetics, 2021, 13, 121.	1.8	13
14	Short-term Exposure to PM2.5 and Epigenetic Aging: A Quasi-Experimental Study. ISEE Conference Abstracts, 2021, 2021, .	0.0	0
15	Effects of short-term personal exposure to air pollution on platelet mitochondrial DNA methylation levels and the potential mitigation by L-arginine supplementation. Journal of Hazardous Materials, 2021, 417, 125963.	6.5	23
16	Short-term exposure to PM2.5 components and renal health: Findings from the Veterans Affairs Normative Aging Study. Journal of Hazardous Materials, 2021, 420, 126557.	6. 5	20
17	Epigenome-wide association study of serum urate reveals insights into urate co-regulation and the SLC2A9 locus. Nature Communications, 2021, 12, 7173.	5 . 8	8
18	Meta-analyses identify DNA methylation associated with kidney function and damage. Nature Communications, 2021, 12, 7174.	5.8	30

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19	Nonsteroidal Antiinflammatory Drugs Modify the Effect of Short-Term Air Pollution on Lung Function. American Journal of Respiratory and Critical Care Medicine, 2020, 201, 374-378.	2.5	12
20	Associations of annual ambient PM2.5 components with DNAm PhenoAge acceleration in elderly men: The Normative Aging Study. Environmental Pollution, 2020, 258, 113690.	3.7	25
21	Association of Neutrophil to Lymphocyte Ratio With Pulmonary Function in a 30-Year Longitudinal Study of US Veterans. JAMA Network Open, 2020, 3, e2010350.	2.8	18
22	Reply to Alexis: Controlled Chamber Studies Showed Protective Effect of Nonsteroidal Antiinflammatory Drugs against Ozone Exposure: The Stage Was Set for Broader Epidemiologic Investigation. American Journal of Respiratory and Critical Care Medicine, 2020, 201, 1583-1584.	2.5	0
23	Epigenetic Link Between Statin Therapy and Type 2 Diabetes. Diabetes Care, 2020, 43, 875-884.	4.3	43
24	Individual species and cumulative mixture relationships of 24-hour urine metal concentrations with DNA methylation age variables in older men. Environmental Research, 2020, 186, 109573.	3.7	16
25	Accelerated epigenetic aging as a risk factor for chronic obstructive pulmonary disease and decreased lung function in two prospective cohort studies. Aging, 2020, 12, 16539-16554.	1.4	13
26	Quantification of the pace of biological aging in humans through a blood test, the DunedinPoAm DNA methylation algorithm. ELife, 2020, 9, .	2.8	268
27	Smoking-Related DNA Methylation is Associated with DNA Methylation Phenotypic Age Acceleration: The Veterans Affairs Normative Aging Study. International Journal of Environmental Research and Public Health, 2019, 16, 2356.	1.2	22
28	Short-term ambient particle radioactivity level and renal function in older men: Insight from the Normative Aging Study. Environment International, 2019, 131, 105018.	4.8	13
29	Comparative validation of an epigenetic mortality risk score with three aging biomarkers for predicting mortality risks among older adult males. International Journal of Epidemiology, 2019, 48, 1958-1971.	0.9	25
30	Impacts of air pollution, temperature, and relative humidity on leukocyte distribution: An epigenetic perspective. Environment International, 2019, 126, 395-405.	4.8	52
31	Oxidative stress and epigenetic mortality risk score: associations with all-cause mortality among elderly people. European Journal of Epidemiology, 2019, 34, 451-462.	2.5	19
32	Serum 25-Hydroxyvitamin D Levels as an Aging Marker: Strong Associations With Age and All-Cause Mortality Independent From Telomere Length, Epigenetic Age Acceleration, and 8-Isoprostane Levels. Journals of Gerontology - Series A Biological Sciences and Medical Sciences, 2019, 74, 121-128.	1.7	25
33	Accelerated DNA methylation age and the use of antihypertensive medication among older adults. Aging, 2018, 10, 3210-3228.	1.4	21
34	Vitamin D status and epigenetic-based mortality risk score: strong independent and joint prediction of all-cause mortality in a population-based cohort study. Clinical Epigenetics, 2018, 10, 84.	1.8	7
35	Leukocyte telomere length and epigenetic-based mortality risk score: associations with all-cause mortality among older adults. Epigenetics, 2018, 13, 846-857.	1.3	21
36	Recurrent tuberculosis in an urban area in China: Relapse or exogenous reinfection?. Tuberculosis, 2017, 103, 97-104.	0.8	39

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37	Associations of self-reported smoking, cotinine levels and epigenetic smoking indicators with oxidative stress among older adults: a population-based study. European Journal of Epidemiology, 2017, 32, 443-456.	2.5	26
38	Tobacco smoking and smoking-related DNA methylation are associated with the development of frailty among older adults. Epigenetics, 2017, 12, 149-156.	1.3	41
39	Associations of Helicobacter pylori infection and chronic atrophic gastritis with accelerated epigenetic ageing in older adults. British Journal of Cancer, 2017, 117, 1211-1214.	2.9	19
40	Community-based active case finding for tuberculosis in rural western China: a cross-sectional study. International Journal of Tuberculosis and Lung Disease, 2017, 21, 1134-1139.	0.6	13
41	The impact of methylation quantitative trait loci (mQTLs) on active smoking-related DNA methylation changes. Clinical Epigenetics, 2017, 9, 87.	1.8	32
42	DNA methylation changes in response to active smoking exposure are associated with leukocyte telomere length among older adults. European Journal of Epidemiology, 2016, 31, 1231-1241.	2.5	11
43	Different definitions of CpG island methylator phenotype and outcomes of colorectal cancer: a systematic review. Clinical Epigenetics, 2016, 8, 25.	1.8	83
44	Tobacco smoking and methylation of genes related to lung cancer development. Oncotarget, 2016, 7, 59017-59028.	0.8	73
45	Relationship of tobacco smoking and smoking-related DNA methylation with epigenetic age acceleration. Oncotarget, 2016, 7, 46878-46889.	0.8	97
46	DNA methylation changes of whole blood cells in response to active smoking exposure in adults: a systematic review of DNA methylation studies. Clinical Epigenetics, 2015, 7, 113.	1.8	330
47	Association ofgyrA/Bmutations and resistance levels to fluoroquinolones in clinical isolates of Mycobacterium tuberculosis. Emerging Microbes and Infections, 2014, 3, 1-5.	3.0	65