James D Stewart

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9768908/publications.pdf

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24 papers 4,464 citations

686830 13 h-index 676716 22 g-index

24 all docs

24 docs citations

times ranked

24

4642 citing authors

#	Article	IF	CITATIONS
1	Characteristics of movers and predictors of residential mobility in the Atherosclerosis Risk in Communities (ARIC) cohort. Health and Place, 2022, 74, 102771.	1.5	4
2	The Associations of Dietary Copper With Cognitive Outcomes. American Journal of Epidemiology, 2022, 191, 1202-1211.	1.6	9
3	Gaseous air pollutants and DNA methylation in a methylome-wide association study of an ethnically and environmentally diverse population of U.S. adults. Environmental Research, 2022, 212, 113360.	3.7	7
4	Outdoor air pollution exposure and inter-relation of global cognitive performance and emotional distress in older women. Environmental Pollution, 2021, 271, 116282.	3.7	13
5	Epigenetically mediated electrocardiographic manifestations of sub-chronic exposures to ambient particulate matter air pollution in the Women's Health Initiative and Atherosclerosis Risk in Communities Study. Environmental Research, 2021, 198, 111211.	3.7	4
6	Long-term particulate matter exposure and bone mineral density in the Women's Health Initiative. ISEE Conference Abstracts, 2021, 2021, .	0.0	0
7	Methylome-wide association study of central adiposity implicates genes involved in immune and endocrine systems. Epigenomics, 2020, 12, 1483-1499.	1.0	6
8	Particulate Matter and Albuminuria, Glomerular Filtration Rate, and Incident CKD. Clinical Journal of the American Society of Nephrology: CJASN, 2020, 15, 311-319.	2.2	61
9	Blood DNA methylation sites predict death risk in a longitudinal study of 12, 300 individuals. Aging, 2020, 12, 14092-14124.	1.4	15
10	Short-term exposure to air pollution and incidence of stroke in the Women's Health Initiative. Environment International, 2019, 132, 105065.	4.8	37
11	Methylome-wide association study provides evidence of particulate matter air pollution-associated DNA methylation. Environment International, 2019, 132, 104723.	4.8	58
12	DNA methylation GrimAge strongly predicts lifespan and healthspan. Aging, 2019, 11, 303-327.	1.4	1,128
13	Air pollution-associated changes in biomarkers of diabetes risk. Environmental Epidemiology, 2019, 3, e059.	1.4	4
14	DNA methylation-based estimator of telomere length. Aging, 2019, 11, 5895-5923.	1.4	198
15	Genome-wide association study and meta-analysis identify loci associated with ventricular and supraventricular ectopy. Scientific Reports, 2018, 8, 5675.	1.6	4
16	GWAS of epigenetic aging rates in blood reveals a critical role for TERT. Nature Communications, 2018, 9, 387.	5.8	151
17	An epigenetic biomarker of aging for lifespan and healthspan. Aging, 2018, 10, 573-591.	1.4	1,552
18	The Association of Long-Term Exposure to Particulate Matter Air Pollution with Brain MRI Findings: The ARIC Study. Environmental Health Perspectives, 2018, 126, 027009.	2.8	76

#	Article	lF	CITATIONS
19	DNA Methylation Signatures of Depressive Symptoms in Middle-aged and Elderly Persons. JAMA Psychiatry, 2018, 75, 949.	6.0	78
20	Epigenetic clock for skin and blood cells applied to Hutchinson Gilford Progeria Syndrome and ex vivo studies. Aging, 2018, 10, 1758-1775.	1.4	406
21	Genetic loci associated with heart rate variability and their effects on cardiac disease risk. Nature Communications, 2017, 8, 15805.	5.8	95
22	Long-term exposure to residential ambient fine and coarse particulate matter and incident hypertension in post-menopausal women. Environment International, 2017, 105, 79-85.	4.8	37
23	Epigenetic clock analysis of diet, exercise, education, and lifestyle factors. Aging, 2017, 9, 419-446.	1.4	521
24	Abstract P261: Genome-wide Association Study of Susceptibility to Particulate Matter-associated Reduced Heart Rate Variability. Circulation, 2016, 133, .	1.6	0