

Ann Moore

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/10232906/publications.pdf>

Version: 2024-02-01

16
papers

1,600
citations

932766

10
h-index

940134

16
g-index

17
all docs

17
docs citations

17
times ranked

3450
citing authors

#	ARTICLE	IF	CITATIONS
1	DNA methylation-based measures of biological age: meta-analysis predicting time to death. <i>Aging</i> , 2016, 8, 1844-1865.	1.4	786
2	Plasma proteomic signature of age in healthy humans. <i>Aging Cell</i> , 2018, 17, e12799.	3.0	325
3	Difference in Muscle Quality over the Adult Life Span and Biological Correlates in the Baltimore Longitudinal Study of Aging. <i>Journal of the American Geriatrics Society</i> , 2014, 62, 230-236.	1.3	123
4	Genome-wide association studies identify 137 genetic loci for DNA methylation biomarkers of aging. <i>Genome Biology</i> , 2021, 22, 194.	3.8	90
5	An integrative cross-omics analysis of DNA methylation sites of glucose and insulin homeostasis. <i>Nature Communications</i> , 2019, 10, 2581.	5.8	62
6	Mitochondrial genetic variation is enriched in G-quadruplex regions that stall DNA synthesis in vitro. <i>Human Molecular Genetics</i> , 2020, 29, 1292-1309.	1.4	36
7	Change in Epigenome-Wide DNA Methylation Over 9 Years and Subsequent Mortality: Results From the InCHIANTI Study. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2016, 71, 1029-1035.	1.7	35
8	Association of Methylation Signals With Incident Coronary Heart Disease in an Epigenome-Wide Assessment of Circulating Tumor Necrosis Factor α . <i>JAMA Cardiology</i> , 2018, 3, 463.	3.0	33
9	DNA methylation signatures reveal that distinct combinations of transcription factors specify human immune cell epigenetic identity. <i>Immunity</i> , 2021, 54, 2465-2480.e5.	6.6	31
10	Influence of cell distribution and diabetes status on the association between mitochondrial DNA copy number and aging phenotypes in the InCHIANTI study. <i>Aging Cell</i> , 2018, 17, e12683.	3.0	26
11	Blood DNA methylation sites predict death risk in a longitudinal study of 12, 300 individuals. <i>Aging</i> , 2020, 12, 14092-14124.	1.4	15
12	Blood DNA Methylation and Aging: A Cross-Sectional Analysis and Longitudinal Validation in the InCHIANTI Study. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2020, 75, 2051-2055.	1.7	14
13	Epigenetic Age Acceleration and Hearing: Observations From the Baltimore Longitudinal Study of Aging. <i>Frontiers in Aging Neuroscience</i> , 2021, 13, 790926.	1.7	9
14	Mitochondrial DNA copy number and heteroplasmy load correlate with skeletal muscle oxidative capacity by P31 MR spectroscopy. <i>Aging Cell</i> , 2021, 20, e13487.	3.0	8
15	Image-based Tissue Distribution Modeling for Skeletal Muscle Quality Characterization. <i>IEEE Transactions on Biomedical Engineering</i> , 2015, 63, 1-1.	2.5	6
16	Prior psychosocial profile and perceived impact of the COVID-19 pandemic: insights from the Baltimore Longitudinal Study of Aging. <i>Aging Clinical and Experimental Research</i> , 2022, 34, 1463-1469.	1.4	1