Ann Moore

List of Publications by Year
in descending order

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