

Epigenetic drift in the aging genome: a ten-year follow-

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Changes in blood DNA methylation and incomplete adjustment for blood cell composition. <i>International Journal of Epidemiology</i> , 2017, 46, 1714-1717.	0.9	3
2	Identification, replication and characterization of epigenetic remodelling in the aging genome: a cross population analysis. <i>Scientific Reports</i> , 2017, 7, 8183.	1.6	27
3	Handling blood cell composition in epigenetic studies on ageing. <i>International Journal of Epidemiology</i> , 2017, 46, 1717-1718.	0.9	6
4	Chronological age prediction based on DNA methylation: Massive parallel sequencing and random forest regression. <i>Forensic Science International: Genetics</i> , 2017, 31, 19-28.	1.6	130
5	A genome-wide association study of cognitive function in Chinese adult twins. <i>Biogerontology</i> , 2017, 18, 811-819.	2.0	18
6	DNA Methylation Changes in Intron 1 of Triggering Receptor Expressed on Myeloid Cell 2 in Japanese Schizophrenia Subjects. <i>Frontiers in Neuroscience</i> , 2017, 11, 275.	1.4	19
7	Implications of DNA Methylation in Parkinson's Disease. <i>Frontiers in Molecular Neuroscience</i> , 2017, 10, 225.	1.4	71
8	Consistency and Variability of DNA Methylation in Women During Puberty, Young Adulthood, and Pregnancy. <i>Genetics & Epigenetics</i> , 2017, 9, 1179237X1772154.	2.5	20
9	Apolipoprotein E DNA methylation and late-life disease. <i>International Journal of Epidemiology</i> , 2018, 47, 899-907.	0.9	22
10	DNA methylation dynamics in aging: how far are we from understanding the mechanisms?. <i>Mechanisms of Ageing and Development</i> , 2018, 174, 3-17.	2.2	135
11	Epigenetic and Transcriptional Variability Shape Phenotypic Plasticity. <i>BioEssays</i> , 2018, 40, 1700148.	1.2	71
12	Back to the future: Epigenetic clock plasticity towards healthy aging. <i>Mechanisms of Ageing and Development</i> , 2018, 174, 18-29.	2.2	71
13	Sex Differences in Aging: Genomic Instability. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2018, 73, 166-174.	1.7	66
14	DNA Methylation and All-Cause Mortality in Middle-Aged and Elderly Danish Twins. <i>Genes</i> , 2018, 9, 78.	1.0	27
15	Epigenetic influences on aging: a longitudinal genome-wide methylation study in old Swedish twins. <i>Epigenetics</i> , 2018, 13, 975-987.	1.3	65
16	DNA methylation 101: what is important to know about DNA methylation and its role in SLE risk and disease heterogeneity. <i>Lupus Science and Medicine</i> , 2018, 5, e000285.	1.1	52
17	DNA Methylation-Based Measures of Biological Aging. , 2018, , 39-64.		16
18	An optimized library for reference-based deconvolution of whole-blood biospecimens assayed using the Illumina HumanMethylationEPIC BeadArray. <i>Genome Biology</i> , 2018, 19, 64.	3.8	245

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19	Epigenetic variability in conversion to psychosis: novel findings from an innovative longitudinal methylomic analysis. <i>Translational Psychiatry</i> , 2018, 8, 93.	2.4	14
20	Defective DNA Methylation/Demethylation Processes Define Aging-Dependent Methylation Patterns. , 2018, , 33-58.		0
21	Genetic background, epigenetic factors and dietary interventions which influence human longevity. <i>Biogerontology</i> , 2019, 20, 605-626.	2.0	32
22	Education and Lifestyle Factors Are Associated with DNA Methylation Clocks in Older African Americans. <i>International Journal of Environmental Research and Public Health</i> , 2019, 16, 3141.	1.2	88
23	The Danish Twin Registry: An Updated Overview. <i>Twin Research and Human Genetics</i> , 2019, 22, 499-507.	0.3	49
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32	Physical Activity, Television Viewing Time, and DNA Methylation in Peripheral Blood. <i>Medicine and Science in Sports and Exercise</i> , 2019, 51, 490-498.	0.2	16
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