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Longitudinal changes in the genetic and environmental influences on the epigenetic clocks across old age: Evidence from two twin cohorts

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24	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,	4.6	49
23	The Danish Twin Registry: An Updated Overview. Twin Research and Human Genetics, 2019, 22, 499-507	2.2	18
22	Should we invest in biological age predictors to treat colorectal cancer in older adults?. <i>European Journal of Surgical Oncology</i> , 2020 , 46, 316-320	3.6	11
21	Association of Blood Chemistry Quantifications of Biological Aging With Disability and Mortality in Older Adults. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2020 , 75, 1671-	16 7 9	17
20	DNA Methylation Clocks and Their Predictive Capacity for Aging Phenotypes and Healthspan. <i>Neuroscience Insights</i> , 2020 , 15, 2633105520942221	3	22
19	Epigenetic measures of ageing predict the prevalence and incidence of leading causes of death and disease burden. <i>Clinical Epigenetics</i> , 2020 , 12, 115	7.7	40
18	A decade of epigenetic change in aging twins: Genetic and environmental contributions to longitudinal DNA methylation. <i>Aging Cell</i> , 2020 , 19, e13197	9.9	11
17	The use of DNA methylation clock in aging research. Experimental Biology and Medicine, 2021, 246, 436-	4 46	3
16	DNA methylation age as a biomarker for cancer. <i>International Journal of Cancer</i> , 2021 , 148, 2652-2663	7.5	6
15	Skewness of X-chromosome inactivation increases with age and varies across birth cohorts in elderly Danish women. <i>Scientific Reports</i> , 2021 , 11, 4326	4.9	6
14	Combined healthy lifestyle score and risk of epigenetic aging: a discordant monozygotic twin study. <i>Aging</i> , 2021 , 13, 14039-14052	5.6	3
13	Epigenetic clocks predict prevalence and incidence of leading causes of death and disease burden.		8
12	Leisure-Time and Occupational Physical Activity Associates Differently with Epigenetic Aging. Medicine and Science in Sports and Exercise, 2021, 53, 487-495	1.2	8
11	Rate of brain aging and are synergistic risk factors for Alzheimeræ disease. <i>Life Science Alliance</i> , 2019 , 2,	5.8	5
10	Epigenetics as a key link between psychosocial stress and aging: concepts, evidence, mechanisms?. <i>Dialogues in Clinical Neuroscience</i> , 2019 , 21, 389-396	5.7	12
9	Epidemiological and genetic overlap among biological aging clocks: New challenges in biogerontology. <i>Ageing Research Reviews</i> , 2021 , 72, 101502	12	2
8	A decade of epigenetic change in aging twins: genetic and environmental contributions to longitudinal DNA methylation.		

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7	The Utility of Twins for Epigenetic Analysis. 2022 , 213-233		Ο
6	Counteracting aged DNA methylation states to combat ageing and age-related diseases. <i>Mechanisms of Ageing and Development</i> , 2022 , 111695	5.6	O
5	Differences in DNA Methylation-Based Age Prediction Within Twin Pairs Discordant for Cancer. 1-9		O
4	Self-reported sleep bruxism in 1990 and 2011 in a nationwide twin cohort: Evidence of trait persistence and genetic liability.		1
3	Epigenetics. 2022 , 585-616		O
2	Epigenetic clocks in neurodegenerative diseases: a systematic review. jnnp-2022-330931		O
1	The relationship of trait-like compassion with epigenetic aging: The population-based prospective Young Finns Study. 14,		O