

## ANNOTATED BIBLIOGRAPHY

**Longitudinal measures of the pace of aging from the Dunedin Study:  
The Pace of Aging, DunedinPoAm, DunedinPACE, and DunedinPACNI  
January 2025, Terrie E. Moffitt**

**Website:** <https://moffittcaspi.trinity.duke.edu/dunedinpace>

### Table of contents

#### Background

**History of research from Pace of Aging to DunedinPACE**

**Six Design Advantages Distinguish PACE From Other aging clocks**

**Generations of measures of biological aging, the 3<sup>rd</sup> generation is: longitudinal**

#### Progress to date

**Ethnic ancestry, does PACE measure ageing in people who are not white?**

**Age, does PACE measure ageing in older adults, and in young people?**

**Risk factors that predict fast aging**

**Predicting clinical endpoints**

**Predicting diagnosed dementia:**

**Responsiveness, the sensitivity of DunedinPACE to detecting change in interventions**

#### Reference list grouped by topics, compiled to 18 January 2025:

#### Groups of Outcomes predicted by DunedinPACE

**Mortality and Frailty**

**Disease Morbidity and Disease Mechanisms**

**Alzheimers, dementias, cognitive decline and brain outcomes**

#### Groups of Predictors of DunedinPACE

**Early-life Risk Factors**

**Health Behaviors, Lifestyles and Psychosocial risks**

**Health Inequalities**

**Mental Health Drugs & Alcohol Abuse**

**Toxic Exposures**

#### Experimental and Intervention Studies of DunedinPACE

#### Methodological Studies of DunedinPACE

#### Reviews of the Literature including DunedinPACE

## BACKGROUND

## HISTORY OF RESEARCH FROM PACE OF AGING TO DUNEDINPACE

Our work aimed to develop a measure that operationalized the definition of aging as: functional decline that (1) simultaneously involves multiple organ systems across the whole body, and (2) is unidirectional, gradual, and progressive. Our Duke/Otago team first modelled the **Pace of**

**Aging** in 2014 using multiple waves of biomarkers assessed in the Dunedin Longitudinal Study when the cohort members were aged 26, 32, 38 (Belsky et al. PNAS 2015). We then updated and improved the Pace of Aging in 2021 by adding an additional wave of data, now covering ages 26, 32, 38, and 45 (Elliot et al. Nature Ageing 2021). The 19 biomarkers repeated every six years indexed variation among individuals in the function of six organ systems: the cardiovascular, metabolic, renal, immune, dental, and pulmonary systems. Dunedin Study participants with faster Pace of Aging scores had more cognitive difficulties, signs of advanced brain aging on MRI, diminished sensory–motor functions, older facial appearance and more pessimistic perceptions of aging.

Unfortunately, however, we realized that few other research teams could adopt our Pace of Aging approach at that time, because they lacked repeated waves of multiple biomarkers across the whole body. To meet this need to export the Pace of Aging to the research community, we translated the Pace of Aging into a DNA methylation version that can be derived from one blood draw. In 2020, we reported how to measure the Pace of Aging by using DNA methylation data, calling the measure **DunedinPoAm** (Belsky et al. eLife 2020).

**DunedinPACE:** In 2022 we updated the DNA methylation version of the Pace of Aging measure to cover the Pace of Aging assessed over ages 26, 32, 38, and 45, naming the new measure DunedinPACE for “Pace of Aging Calculated in the Epigenome.” The details of DunedinPACE are published in a scientific article (Belsky et al. 2022 eLife, link [here](#)). DunedinPACE is an epigenetic biomarker that measures the human pace of aging. It is designed to function as a speedometer for aging, offering a single-timepoint measurement of how fast a person is now aging. DunedinPACE is designed to function as a speedometer for aging, offering a single-timepoint measurement of how fast a person’s body is now aging. We exported DunedinPACE to five additional cohorts, and found that it showed very high test-retest reliability, predicted morbidity, disability, and mortality, and indicated faster aging in adults who survived childhood adversity. The code to compute DunedinPACE in DNA methylation data from Illumina 450k or EPIC1 or EPIC2 arrays is available open-access on GitHub, [here](#). The code is also included in [BioLearn](#). (NOTE: because the Illumina company who provides DNA methylation arrays has updated from their 450k to EPIC1 to EPIC2, we have updated the Github algorithm. The algorithm automatically works with data created from any of those three Illumina platforms.)

**DunedinPACNI:** Despite the success of DunedinPACE in cohorts that have DNA methylation data, we realized that many important studies of aging have not invested in costly DNA methylation, but instead have invested in collecting brain MRI data. As a result, in 2024 we translated the Pace of Aging into a brain-MRI version that can be derived from one MRI brain scan image. We call this measure **DunedinPACNI** for “Pace of Aging Calculated in a Neuro-Image” (Whitman et al. 2025). It has very high test-retest reliability. We exported this measure to the Alzheimer’s Disease Neuroimaging Initiative, the UK Biobank, and the BrainLAT neuroimaging datasets and found that faster DunedinPACNI predicted participants’ cognitive impairment, accelerated brain atrophy, and conversion from mild cognitive impairment to diagnosed dementia. Underscoring close links between longitudinal aging of the body and brain, faster DunedinPACNI also predicted physical frailty, poor health, future chronic diseases, and mortality in older adults. The code to computer DunedinPACNI is here: XXXX.

## Six Design Advantages Distinguish this Measure from Others

First, DunedinPACE was derived from 19 biomarkers ascertained longitudinally in a cohort over 2 decades at ages 26, 32, 38, and 45, comprising 69,715 data points. In this longitudinal design the participants were tested four times, each 5 years apart. This spacing allowed us to track unidirectional (i.e., downhill) decline that is correlated across organ systems as age advances gradually, long-term. This is an advantage because DunedinPACE has no noise from short term illnesses or infections, such as an infected tooth or a viral flu, that could temporarily elevate or spike a biomarker during a spell of illness. Most other bio-age measures are derived from only one cross-section of data and thus they inadvertently contain some noise from spells of sickness.

Second, DunedinPACE was derived on a cohort of research participants who were all born the same year. To date this design feature is unique in the world of aging measures. This is an advantage because DunedinPACE has no noise from historical differences in exposure to factors that in multi-age samples alter the participants' body or their epigenome. For example, older generations had more exposure to toxins like cigarette smoke and leaded gasoline, fewer years of education, and few fruits and vegetables in winter. Younger generations had childhood vaccinations, antibiotics, air-conditioning, anti-inflammatory medications, and oral contraception. These generational factors can alter the epigenome, but they are not relevant to aging. Other bio-age measures that were derived on people who have different birth years inadvertently contain some of this exposure noise.

Third, DunedinPACE was derived in a cohort of adults studied up to age 45. This is an advantage because DunedinPACE has no noise from late-life diseases such as diabetes, heart disease, kidney disease, dementia, or cancers, because these diseases had not yet onset in the Dunedin cohort when we constructed DunedinPACE. We aimed for a measure that reflects aging and is not merely a repackaging of disease. Other bio-age measures are typically derived from samples in which many of the older members already have chronic diseases that have altered their body and epigenome, so those measures can reflect disease more than aging itself.

Fourth, DunedinPACE was derived in a midlife cohort for which attrition by death has been minimal; under 3%. This is an advantage because of so-called "survivor bias." Other bio-age measures are typically derived from samples in which the many of the older members are missing due to death. In multi-age samples, the youngest participants can be sampled from all people born their year, but the older participants can only be sampled from the subset of those left alive. In mixed-age samples with attrition and in studies that begin in midlife or later, the fastest agers will have already died, which reduces the potential range of scores on any bio-age measure. Thus, in other bio-age measures, the young versus old participants differed from each other on some noise factors that are not relevant for aging.

Fifth, DunedinPACE was derived from a pool of DNA methylation probe sites across the epigenome that were pre-selected to have strong test-retest reliability. Unreliable probe sites were excluded from DunedinPACE. We showed that most methylation probe sites are indeed unreliable, which adds noise to DNAm bio-age measures. Other bio-age measures sometimes offset this noise after the fact using statistical procedures, such as deriving principal factors. DunedinPACE builds reliability in.

Sixth, because we intended it to be responsive to anti-aging interventions, unlike other aging clocks DunedinPACE was trained on change over time. Evidence is accumulating that

DunedinPACE Among aging clocks, DunedinPACE is the most sensitive to intervention to date, slows most in intervention trials, and is the most consistently responsive across studies.

**THE THIRD GENERATION IS LONGITUDINAL.** DunedinPACE is sometimes referred to as a “third generation” among clocks, because earlier generations of clocks were not derived in longitudinal data to assess actual physiological decline in healthy people who are all the same chronological age.

## PROGRESS TO DATE

**More than 65 large epidemiological cohort studies in 17 countries are now using DunedinPACE**, including, in alphabetical order: Add Health, The Alzheimer disease neuroimaging initiative (ADNI), AHAB, ALSPAC UK, ASPREE Study Australia, Austrian Stroke Prevention Study, BeCOME Germany, Baltimore Longitudinal Study of Aging (BLSA), BrainLAT, CALERIE, Canadian longitudinal study on aging (CLSA), CARDIA, Cebu Longitudinal Health and Nutrition Survey (CLHNS Philippines), Child Health and Development Study, Chinese National Twin Registry, COBRA, Dutch Hunger Winter Study, E-Risk Longitudinal Twin Study UK, FACHS (African American families), FinnTwin, Finnish Twin Study on Aging FITSA, Young Finn Study, Framingham Offspring Cohort, Fragile Families, Future of Families and Child Wellbeing (FFCW), Generation Scotland, German Socioeconomic Panel Study (SOEP-Gene), Grady Trauma Project, HANDLE (NIA’s intramural study), Health and Retirement Study (HRS), InChianti, the Korean Genome and Epidemiology Study (KoGES), KORA (Cooperative Health Research in the Region Augsburg study), Leiden Longevity Study, Lothian Birth cohort, Melbourne Collaborative Cohort Study Australia, MESA, MIDUS, United States National Health and Nutrition Examination Survey (NHANESIII), National Heart, Lung, and Blood Institute Growth and Health Study (NGHS, 1992), Netherlands Twin Registry, NHANES, Normative Aging Study, Northern Finland 1966 birth cohort, Norwegian MOBA, Penn State Child Health Study, REWARD Study in Wisconsin, Rotterdam Study, Saint Jude Lifetime Cohort of Cancer Survivors (SJLIFE), Sister Study, SOL-INCA and HCHS-SOL, SOEPG Germany, Strong Heart Study of Native American Indians, Swedish Adoption/Twin Study of Aging, Swiss Family Study, Taiwanese Biobank, The Texas Twin Study, TILDA Ireland, Thinking and Living with Cancer (TLC) Study, TwinLife, UK Household Longitudinal Study, UK Understanding Society, Veterans Administration Post-Deployment Mental Health Study (PDMH), Women and their Children’s Health in Southeast Louisiana, Women’s Health Initiative, plus others.

## STUDIES VALIDATING DUNEDINPACE AS A USEFUL MEASURE OF WHOLE-BODY AGING:

**Other research teams have published more than 300 reports** of new findings from these data sets since the measure became available to them in 2021 (these reports are not from our team).

**We ourselves have published more than 35 reports** with findings using our measures of the pace of aging. As an example, we reported that faster DunedinPACE distinguishes long-term cannabis users from recreational users (Meier et al. 2022, *Lancet Healthy Longevity*). As another example, people with more education have slower DunedinPACE, even after controls for genetic confounds (Sugden et al. 2023, *J of Gerontology*).

**Replication and robustness.** Our publications have generally replicated findings by using one or more of the data sets that now include DunedinPACE; we often report multi-study

publications. Our paper announcing DunedinPACE reported validation in 5 data sets (Belsky et al 2022, *eLife*). As another example, we reported that faster DunedinPACE distinguishes Alzheimer dementia patients in **2 cohorts**, ADNI and the Framingham Heart Study (Sugden et al. 2022, *Neurology*). We also reported that people with more education have slower DunedinPACE, after controls for genetic confounds, in **5 cohorts** (Sugden et al. 2023, *J of Gerontology*). Other replication papers show that DunedinPACE relates to MRI-measured brain structure in **3 cohorts** (Whitman et al. 2023 *Neurobiology of Aging*) and to schizophrenia in **5 case-control studies** (Caspi et al. 2023 *Biological Psychiatry*).

**Comparison to clocks:** Overall, findings have shown that DunedinPACE routinely outperforms the Horvath, Hannum, and PhenoAGE clocks, and typically performs on par with the GrimAGE clock. DunedinPACE outperformed the GrimAGE clock on some analyses of cognitive decline and Alzheimers dementia, and in studies of responsiveness to interventions, but these comparisons are early days yet.

**Ethnic ancestry:** DunedinPACE was originally trained on white New Zealanders, but it is being reported on groups of other ancestries. Yin (2023) reported in the Taiwanese Biobank that DunedinPACE is faster in **Han Chinese** who have multimorbid physical illnesses. Kim (2024) reported that DunedinPACE is faster in the Korean Genome and Epidemiology Study (KoGES) in **Korean** participants who have chronic disease and poor health lifestyle factors. Boyer et al (2023) reported DunedinPACE findings in **Native American Indians**. Overall, published findings in **African Americans** are looking similar to those in Whites, as shown in the FACHS, HANDLE, Future of Families and Child Wellbeing (FFCW), MIDUS, National Heart, Lung, and Blood Institute Growth and Health Study (NGHS 1992), and HRS data sets, as well as the Veterans Administration Post-Deployment Mental Health Study (PDMH), which is 50% African American. Krieger et al. studied DunedinPoAm in Black, Hispanic, and white ethnicity with respect to racialized and economic injustice in MESA. Gibbs et al. reported on DunedinPACE in **Pasifika**

**Age, children:** DunedinPACE was originally trained on people age 45, but overall, published findings show DunedinPACE is relevant for studies of children and adolescents: E-Risk, ALSPAC, FinnTwin, PennState Child Health Study, Texas Twins, Quebec Longitudinal Study of Child Development, Future of Families and Child Wellbeing (FFCW), and others. More data are needed to understand interpretation of scores in children. (NOTE: some child studies used buccal/saliva as the source of DNA, whereas DunedinPACE was trained on venous blood as the source of DNA. The implications of this are not yet clear.)

**Age, older adults:** DunedinPACE was trained on people age 45, but it is proving relevant for studies of older adults: The Alzheimer disease neuroimaging initiative (ADNI), NIA's Baltimore Longitudinal Study on Aging (BLSA), Berlin Study of Aging, Framingham Heart Study, Healthy Aging in Neighborhoods of Diversity Across the Life Span study (HANDLE), Health and Retirement Study (HRS), Multi-Ethnic Study of Atherosclerosis (MESA), Normative Aging Study, The Irish Longitudinal Study on Ageing (TILDA), UK Understanding Society, Finnish Twin Study on Aging (FITSA), Canadian longitudinal study on aging (CLSA), 1958 British Birth cohort, Scottish Lothian birth Cohorts, Swedish Adoption/Twin Study of Aging (SATSA), Rotterdam Aging Study, Leiden Longevity Study, Women's Health Initiative, and others.

**Predicting clinical endpoints:** Faster DunedinPACE has been shown to statistically predict endpoint indicators including functional and cognitive decline, frailty, disease multimorbidity, and mortality (age at death) in AHAB, BLSA, Framingham, Generation Scotland, HANDLE, HRS, Melbourne Collaborative Cohort Study, MESA, NAM, UK Understanding Society, FITSA, Taiwanese Biobank, Rotterdam Study, Leiden Longevity Study, Lothian birth cohort, Women's Health Initiative, and others.

**Predicting diagnosed dementia:** Faster DunedinPACE has been reported to predict ADRD and Mild Cognitive Impairment in ADNI and Framingham (Savin et al, 2024; Sugden et al, 2022), and cognitive decline (Elliot et al. 2021; Reed et al. 2022). Another study did not find this (Schäfer Hackenhaar, et al. 2023).

**Risk factors that predict fast aging:** DunedinPACE has been shown to be accelerated in people who experienced psycho-social adversity and deprivation in AddHealth, ALSPAC, BLSA (Baltimore Longitudinal Study of Aging), E-Risk Longitudinal Twin Study UK, FACHS (African American families study), Future of Families and Child Wellbeing (FFCW), Generation Scotland, HANDLE (NIA's intramural study), HRS, MESA, NHANES, Normative Aging Study, the Sister Study, Swiss Family Study, The Texas Twin Study, TILDA (Ireland), UK Understanding Society, Canadian longitudinal study on aging (CLSA), Grady Trauma Project, Taiwanese Biobank, Veterans Administration Post-Deployment Mental Health Study, and five schizophrenia case-control studies, plus other studies.

**Responsiveness, the sensitivity of DunedinPACE to detecting change in interventions:** Sehgal et al (in press) re-analysed 51 longevity intervention trials in humans, and systematically tested 16 DNAm clocks across the 51 intervention trials. They concluded that among 16 aging clocks, DunedinPACE had the most evidence for sensitivity to intervention so far. DunedinPACE had the largest mean decrease in anti-aging intervention trails, and DunedinPACE was the most consistently responsive, significantly decreasing in 16 interventions and increasing in only 1. As an example, DunedinPACE was reported to slow in response to caloric restriction in the CALERIE trial. DunedinPACE has also slowed after successful treatment for insomnia. DunedinPACE was sensitive to a drug intervention in ClockBASE. It has also been shown to accelerate under conditions of psychological stress in FACHS, under conditions of bio-stress in three clinical datasets reported by Poganik et al. 2023, and in patients who are badly burned (Sullivan et al. 2025). This set of findings is potentially important because it shows that DunedinPACE can be sensitive to detecting change, and that it can be repeated as a before-after measure on a time-span of weeks to quantify short-term change in the pace of biological aging. But this is early days and much more research is needed to confirm.

## GROUPED REFERENCE LISTS:

Literature reporting findings up to the date of this report for the DunedinPACE Pace of Aging measure (or prior version, PoAm):

The name of the dataset is below each citation.

### PUBLICATIONS ON FRAILTY and MORTALITY

Balachandran, A., Pei, H., Beard, J., Caspi, A., Cohen, A., Domingue, B. W., ... & Belsky, D. W. (2024). Pace of Aging in older adults matters for healthspan and lifespan. *medRxiv*. ELSA, HRS

Belsky, D. W., Caspi, A., Arseneault, L., Baccarelli, A., Corcoran, D. L., Gao, X., Hannon, E., Harrington, H. L., Rasmussen, L. J., Houts, R., Huffman, K., Kraus, W. E., Kwon, D., Mill, J., Pieper, C. F., Prinz, J. A., Poulton, R., Schwartz, J., Sugden, K., . . . Moffitt, T. E. (2020). Quantification of the pace of biological aging in humans through a blood test, the DunedinPoAm DNA methylation algorithm. *eLife*, 9. <https://doi.org/10.7554/eLife.54870>  
Normative Aging Study, Framingham

Belsky, D. W., Caspi, A., Corcoran, D. L., Sugden, K., Poulton, R., Arseneault, L., Baccarelli, A., Chamarti, K., Gao, X., Hannon, E., Harrington, H. L., Houts, R., Kothari, M., Kwon, D., Mill, J., Schwartz, J., Vokonas, P., Wang, C., Williams, B. S., & Moffitt, T. E. (2022). DunedinPACE, a DNA methylation biomarker of the pace of aging. *eLife*, 11. <https://doi.org/10.7554/eLife.73420>  
Normative Aging Study, Framingham

Bernabeu, E., McCartney, D. L., Gadd, D. A., Hillary, R. F., Lu, A. T., Murphy, L., Wrobel, N., Campbell, A., Harris, S. E., Liewald, D., Hayward, C., Sudlow, C., Cox, S. R., Evans, K. L., Horvath, S., McIntosh, A. M., Robinson, M. R., Vallejos, C. A., & Marioni, R. E. (2023). Refining epigenetic prediction of chronological and biological age. *Genome Med*, 15(1), 12. <https://doi.org/10.1186/s13073-023-01161-y>  
Lothian Birth Cohort, Generation Scotland, Framingham Study

Bourassa, K. J., Anderson, L., Woolson, S., Dennis, P. A., Garrett, M. E., Hair, L., Dennis, M., Sugden, K., Williams, B. S., Houts, R., VA Mid Atlantic MIRECC Workgroup, Calhoun, P. S., Naylor, J., Ashley-Koch, A. E., Beckham, J. C., Caspi, A., Taylor, G. A., Hall, K. S., Moffitt, T. E., & Kimbrel, N. A. (2024). Epigenetic aging and prospective morbidity and mortality among U.S. veterans.  
Veterans Administration Post-Deployment Mental Health Study (PDMH).

Cribb, L., Hodge, A. M., Yu, C., Li, S. X., English, D. R., Makalic, E., Southey, M. C., Milne, R. L., Giles, G. G., & Dugué, P. A. (2022). Inflammation and Epigenetic Aging Are Largely Independent Markers of Biological Aging and Mortality. *J Gerontol A Biol Sci Med Sci*, 77(12), 2378-2386. <https://doi.org/10.1093/gerona/glac147>  
Melbourne Collaborative Cohort Study

**Crimmins:** Crimmins, E. M., Klopack, E. T., & Kim, J. K. (2024). Generations of epigenetic clocks and their links to **socioeconomic status** in the Health and Retirement Study. *Epigenomics*, 1–12. <https://doi.org/10.1080/17501911.2024.2373682>  
The Health and Retirement Study (HRS)

Daredia, Saher, Dennis Khodasevich, Nicole Gladish, Hanyang Shen, Jamaji C. Nwanaji-Enwerem, Anne K. Bozack, Belinda L. Needham, David H. Rehkoppf, Julianna Deardorff, Andres Cardenas. Timing of Menarche and Menopause and Epigenetic Aging among U.S. Adults: Results from the National Health and Nutrition Examination Survey 1999-2002  
medRxiv 2024.12.19.24319271; doi: <https://doi.org/10.1101/2024.12.19.24319271>  
NHANES

Faul, J. D., Kim, J. K., Levine, M. E., Thyagarajan, B., Weir, D. R., & Crimmins, E. M. (2023). Epigenetic-based age acceleration in a representative sample of older Americans: Associations with aging-related morbidity and mortality. *Proc Natl Acad Sci U S A*, 120(9), e2215840120. <https://doi.org/10.1073/pnas.2215840120>  
HRS

Föhr, T., Waller, K., Viljanen, A., Rantanen, T., Kaprio, J., Ollikainen, M., & Sillanpää, E. (2023). Mortality Associations With DNA Methylation-Based Biological Aging and Physical Functioning Measures Across a 20-Year Follow-up Period. *J Gerontol A Biol Sci Med Sci*, 78(8), 1489-1496. <https://doi.org/10.1093/gerona/glad026>  
FITSA Finnish Twin Study on Aging

Fong, Sheng, Kirill A. Denisov, Brian K. Kennedy, Jan Gruber (2024). LinAge2: Providing actionable insights and benchmarking with epigenetic clocks. BioRxiv.  
doi: <https://doi.org/10.1101/2024.12.23.24319587>  
NHANES

Guida, J. L., Hyun, G., Belsky, D. W., Armstrong, G. T., Ehrhardt, M. J., Hudson, M. M., Green, P. A., Robison, L. L., Streck, B. P., Tonorezos, E. S., Yasui, Y., Wilson, C. L., Wang, Z., & Ness, K. K. (2024). Associations of seven measures of biological age acceleration with frailty and all-cause mortality among adult survivors of childhood cancer in the St. Jude Lifetime Cohort. *Nature Cancer*. <https://doi.org/10.1038/s43018-024-00745-w>  
St Jude LIFE cohort, NHANESIII

Klopack, Eric T., Eileen M. Crimmins (2024). Epigenetic Aging Helps Explain Differential Resilience in Older Adults. *Demography* 61 (4): 1023–1041.  
<https://doi.org/10.1215/00703370-11466635>  
HRS

Kuiper, L. M., Polinder-Bos, H. A., Bizzarri, D., Vojinovic, D., Vallerga, C. L., Beekman, M., Dollé, E. T., Ghanbari, M., Voortman, T., Reinders, M. J. T., Verschuren, W. M. M., Slagboom, P. E., van den Akker, E. B., & van Meurs, J. B. J. (2023). Epigenetic and Metabolomic Biomarkers for Biological Age: A Comparative Analysis of Mortality and Frailty Risk. *J Gerontol A Biol Sci Med Sci*, 78(10), 1753-1762.  
<https://doi.org/10.1093/gerona/glad137>  
Rotterdam Study, Leiden Longevity Study

Kuo, Pei-Lun, Ann Zenobia Moore, Toshiko Tanaka, Daniel W Belsky, Ake Tzu-Hui Lu, Steve Horvath, Stefania Bandinelli, Luigi Ferrucci (2024) Longitudinal changes in epigenetic clocks predict survival in the InCHIANTI cohort, BioRxiv, <https://doi.org/10.1101/2024.09.13.24313620>  
Baltimore Longitudinal Study of Aging

Li, D. L., Hodge, A. M., Southey, M. C., Giles, G. G., Milne, R. L., & Dugué, P.-A. (2024). Self-rated health, epigenetic ageing, and long-term mortality in older Australians. *Geroscience*. <https://doi.org/10.1007/s11357-024-01211-2>  
Melbourne Collaborative Cohort Study, MCCS

Mak, J. K. L., Karlsson, I. K., Tang, B., Wang, Y., Pedersen, N. L., Hägg, S., Jylhävä, J., & Reynolds, C. A. (2023). Temporal dynamics of epigenetic aging and frailty from midlife to old age. *J Gerontol A Biol Sci Med Sci*. <https://doi.org/10.1093/gerona/glad251>  
Swedish Adoption and twin Study of Aging (SATSA)

Mendy, A., & Mersha, T. B. (2024). Epigenetic age acceleration and mortality risk prediction in US adults. *medRxiv*.

NHANES

Phyo, A. Z. Z., Fransquet, P. D., Wrigglesworth, J., Woods, R. L., Espinoza, S. E., & Ryan, J. (2024). Sex differences in biological aging and the association with clinical measures in older adults. *Geroscience*, 46(2), 1775-1788. <https://doi.org/10.1007/s11357-023-00941-z>  
ASPREE

Phyo, A. Z. Z., Espinoza, S. E., Murray, A. M., Fransquet, P. D., Wrigglesworth, J., Woods, R. L., & Ryan, J. (2024). Epigenetic age acceleration and the risk of frailty, and persistent activities of daily living (ADL) disability. *Age and Ageing*, 53(6), afae127.  
ASPREE

Rosko, A. E., Elsaied, M. I., Woyach, J., Islam, N., Lepola, N., Urrutia, J., ... & Burd, C. E. (2024). Determining the relationship of p16INK4a and additional molecular markers of aging with clinical frailty in hematologic malignancy. *Journal of Cancer Survivorship*, 1-11.

Sabbatinelli, J., Giuliani, A., Kwiatkowska, K. M., Matacchione, G., Belloni, A., Ramini, D., ... & Bronte, G. (2024). DNA Methylation-derived biological age and long-term mortality risk in subjects with type 2 diabetes. *Cardiovascular diabetology*, 23(1), 250.

Sehgal, R., Meer, M., Shadyab, A. H., Casanova, R., Manson, J. E., Bhatti, P., Crimmins, E. M., Assimes, T. L., Whitsel, E. A., Higgins-Chen, A. T., & Levine, M. (2023). Systems Age: A single blood methylation test to quantify aging heterogeneity across 11 physiological systems. *bioRxiv*. <https://doi.org/10.1101/2023.07.13.548904>  
Women's Health Initiative

Verschoor, C. P., Lin, D. T. S., Kobor, M. S., Mian, O., Ma, J., Pare, G., & Ybazeta, G. (2021). Epigenetic age is associated with baseline and 3-year change in frailty in the Canadian Longitudinal Study on Aging. *Clin Epigenetics*, 13(1), 163.  
<https://doi.org/10.1186/s13148-021-01150-1>  
Canadian longitudinal study on aging (CLSA)

Whitman E.T., Elliott, M.L., Knodt, A.R., Abraham, W.C., Anderson, T.J., Cutfield, N., Hogan, S., Ireland, D., Melzer, T.R., Ramrakha, S., Sugden, K., Theodore, R., Williams, B.S., Caspi, A., Moffitt, T.E., & Hariri, A.R. for the Alzheimer's Disease Neuroimaging Initiative. (Submitted2024). An estimate of the longitudinal pace of aging from a single brain scan predicts dementia conversion, morbidity, and mortality.

ADNI, UK BIOBANK, BRAINLAT

Ying, Kejun, Seth Paulson, Alec Eames, Alexander Tyshkovskiy, Siyuan Li, Martin Perez-Guevara, Mehrnoosh Emamifar, Maximiliano Casas Martínez, Dayoon Kwon, Anna Kosheleva, Michael P. Snyder, Dane Gobel, Chiara Herzog, Jesse R. Poganik, Biomarker of Aging Consortium, Mahdi Moqri, Vadim N. Gladyshev. (2024 preprint) A Unified Framework for Systematic Curation and Evaluation of Aging Biomarkers.

<https://www.researchsquare.com/article/rs-4481437/v1.pdf>

Normative Aging Study, MassGeneral Brigham dataset

Zhang, H., Hao, D., Gong, Y., Xu, Y., Ding, C., Wang, J., ... & Li, X. Associations of Twelve DNA Methylation Signatures of Aging with Mortality. Available at SSRN 4979833.  
NHANES

### PUBLICATIONS ON DISEASE MORBIDITY AND DISEASE MECHANISMS

Agbor, F., Stoudmire, T., Goodin, B., & Aroke, E. (2024). The Pace of Biological Aging Mediates the Relationship between Internalized Chronic Pain Stigma and Pain-related Disability. *The Journal of Pain*, 25(4), 43.

Arge, L. A., Lee, Y., Skåra, K. H., Myrskylä, M., Ramlau-Hansen, C. H., Håberg, S. E., & Magnus, M. C. (2024). Epigenetic aging and fecundability: the Norwegian Mother, Father and Child Cohort Study. *Human Reproduction*, deae242.  
Norwegian MOBA

Aroke, E. N., Srinivasasainagendra, V., Kottae, P., Quinn, T. L., Wiggins, A. M., Hobson, J., Kinnie, K., Stoudmire, T., Tiwari, H. K., & Goodin, B. R. (2023). The Pace of Biological Aging Predicts Nonspecific Chronic Low Back Pain Severity. *J Pain*.  
<https://doi.org/10.1016/j.jpain.2023.10.018>

Aroke, Edwin N, Jai Ganesh Nagidi, Vinodh Srinivasasainagendra, Tammie L Quinn, Fiona BAT Agbor, Kiari R Kinnie, Hemant K Tiwari & Burel R Goodin (2024) The Pace of Biological Aging Partially Explains the Relationship Between Socioeconomic Status and Chronic Low Back Pain Outcomes, *Journal of Pain Research*, , 4317-4329, DOI: 10.2147/JPR.S481452

Arzu, J. L., Kelsey, K. T., Papandonatos, G. D., Cecil, K. M., Chen, A., Langevin, S. M., ... & Braun, J. M. (2024). Associations of epigenetic age acceleration at birth and age 12 years with adolescent cardiometabolic risk: the HOME study. *Clinical Epigenetics*, 16(1), 163.

### The HOME Study

- Baldelli, L., Pirazzini, C., Sambati, L., Ravaioli, F., Gentilini, D., Calandra-Buonaura, G., Guaraldi, P., Franceschi, C., Cortelli, P., Garagnani, P., Bacalini, M. G., & Provini, F. (2023). Epigenetic clocks suggest accelerated aging in patients with isolated REM Sleep Behavior Disorder. *NPJ Parkinsons Dis*, 9(1), 48. <https://doi.org/10.1038/s41531-023-00492-2>
- Bourassa, K. J., Anderson, L., Woolson, S., Dennis, P. A., Garrett, M. E., Hair, L., Dennis, M., Sugden, K., Williams, B. S., Houts, R., VA Mid Atlantic MIRECC Workgroup, Calhoun, P. S., Naylor, J., Ashley-Koch, A. E., Beckham, J. C., Caspi, A., Taylor, G. A., Hall, K. S., Moffitt, T. E., & Kimbrel, N. A. (2024). Epigenetic aging and prospective morbidity and mortality among U.S. veterans. Veterans Administration Post-Deployment Mental Health Study (PDMH).
- Casas, Sandra R., Nuria, M., Nuria, O., Lidia, P., Tamara, G., Oriol, S., Àlvar, A., & Rosa, F. (2023). Is the association between the severity of airflow limitation and DNA Methylation similar in blood and lung tissue in patients with COPD? *European Respiratory Journal*, 62(suppl 67), PA5194. <https://doi.org/10.1183/13993003.congress-2023.PA5194>
- Chang, X. Y., & Lin, W. Y. (2023). Epigenetic age acceleration mediates the association between smoking and diabetes-related outcomes. *Clin Epigenetics*, 15(1), 94. <https://doi.org/10.1186/s13148-023-01512-x>
- Cribb, L., Hodge, A. M., Yu, C., Li, S. X., English, D. R., Makalic, E., Southey, M. C., Milne, R. L., Giles, G. G., & Dugué, P. A. (2022). Inflammation and Epigenetic Aging Are Largely Independent Markers of Biological Aging and Mortality. *J Gerontol A Biol Sci Med Sci*, 77(12), 2378-2386. <https://doi.org/10.1093/gerona/glac147>  
Melbourne Collaborative Cohort Study
- Demuth, Ilja , Valentin Max Vetter, Jan Homann , Vera Regitz-Zagrosek, Denis Gerstorf , Christina M. Lill, Lars Bertram. DunedinPACE Predicts Incident Metabolic Syndrome: Cross-sectional and Longitudinal Data from the Berlin Aging Study II (BASE-II) MedRxiv posted December 16, 2024.;<https://doi.org/10.1101/2024.12.16.24319083>doi: Berlin Aging Study II (BASE II)
- Etzel, L., Hastings, W. J., Hall, M. A., Heim, C. M., Meaney, M. J., Noll, J. G., O'Donnell, K. J., Pokhvisneva, I., Rose, E. J., Schreier, H. M. C., Shenk, C. E., & Shalev, I. (2022). Obesity and accelerated epigenetic aging in a high-risk cohort of children. *Scientific Reports*, 12(1), 8328. <https://doi.org/10.1038/s41598-022-11562-5>
- Faul, J. D., Kim, J. K., Levine, M. E., Thyagarajan, B., Weir, D. R., & Crimmins, E. M. (2023). Epigenetic-based age acceleration in a representative sample of older Americans: Associations with aging-related morbidity and mortality. *Proc Natl Acad Sci U S A*, 120(9), e2215840120. <https://doi.org/10.1073/pnas.2215840120>  
HRS
- Föhr, T., Hendrix, A., Kankaanpää, A., Laakkonen, E. K., Kujala, U., Pietiläinen, K. H., Lehtimäki, T., Kähönen, M., Raitakari, O., Wang, X., Kaprio, J., Ollikainen, M., & Sillanpää, E. (2024). Metabolic syndrome and epigenetic aging: a twin study. *Int J Obes (Lond)*. <https://doi.org/10.1038/s41366-024-01466-x>

### Finnish Twins

Freij, Khalid W., Fiona B.A.T. Agbor, Kiari R. Kinnie, Vinodh Srinivasasainagendra, Tammie L. Quinn, Hemant K. Tiwari, Robert E. Sorge, Burel R. Goodin, Edwin N. Aroke, (2024) The pace of biological aging significantly mediates the relationship between internalized stigma of chronic pain and chronic low back pain severity among non-hispanic black but not non-hispanic white adults, *Neurobiology of Pain*, Volume 16, 2024, 100170, ISSN 2452-073X, <https://doi.org/10.1016/j.ynpai.2024.100170>.

Garrett, Melanie E. MS<sup>1</sup>, Brandon Le, PhD<sup>1</sup>, Kyle J. Bourassa, PhD<sup>2,3</sup>, Michelle F. Dennis, BA<sup>2,4</sup>, Daniel Hatch, PhD<sup>5</sup>, Qing Yang, PhD<sup>5</sup>, Paula Tanabe, PhD<sup>5</sup>, Nirmish Shah, MD<sup>6</sup>, Faith S. Luyster, PhD<sup>7</sup>, Charity Oyedeleji, MD<sup>6,8</sup>, John J. Strouse, MD, PhD<sup>6,8</sup>, Harvey J. Cohen, MD<sup>3</sup>, Nathan A. Kimbrel, PhD<sup>2,4</sup>, Jean C. Beckham, PhD<sup>2,4</sup>, Mitchell R. Knisely, PhD<sup>5</sup>, Marilyn J. Telen, MD<sup>6</sup>, Allison E. Ashley-Koch, PhD<sup>1\*</sup>, for the VA Mid-Atlantic MIRECC Workgroup. (2024). Black Americans with sickle cell disease (SCD) demonstrate accelerated epigenetic pace of aging compared to Black Americans without SCD. *Journals of Gerontology, Series A: Medical Sciences*.

Gehle, S. C., Kleissler, D., Heiling, H., Deal, A., Xu, Z., Ayer Miller, V. L., Taylor, J. A., & Smitherman, A. B. (2023). Accelerated epigenetic aging and myopenia in young adult cancer survivors. *Cancer Med*, 12(11), 12149-12160. <https://doi.org/10.1002/cam4.5908>

Gibbs, S., Jones, G., Cameron, V., Fakahau, P., Frampton, C., Earle, N., ... & Fa'atoese, A. (2024). Epigenetic Measures of 'Biological Ageing' Associated With Cardiovascular Risk in Non-Pacific Populations do not Predict Cardiovascular Risk in a Cohort of Samoans, Tongans, and Fijians Living in Aotearoa New Zealand, except for DunedinPACE. *Heart, Lung and Circulation*, 33, S59-S60.

Pasifika Heart Study

Hillary, R. F., Stevenson, A. J., McCartney, D. L., Campbell, A., Walker, R. M., Howard, D. M., Ritchie, C. W., Horvath, S., Hayward, C., McIntosh, A. M., Porteous, D. J., Deary, I. J., Evans, K. L., & Marioni, R. E. (2020). Epigenetic measures of ageing predict the prevalence and incidence of leading causes of death and disease burden. *Clin Epigenetics*, 12(1), 115. <https://doi.org/10.1186/s13148-020-00905-6>

Generation Scotland: Scottish Family Health Study

Ji, J., Sun, C. L., Cohen, H. J., Freedman, R. A., Chapman, A. E., Klepin, H. D., ... & Sedrak, M. S. (2024). SIOG2024-2-P-038 Blood-based DNA methylation measures of epigenetic age acceleration in older women with and without breast cancer. *Journal of Geriatric Oncology*, 15(7), 101895.

Kim DJ, Kang JH, Kim J-W, Kim Sb, Lee YK, Cheon MJ and Lee B-C (2024) Assessing the utility of epigenetic clocks for health prediction in South Korean. *Front. Aging* 5:1493406. doi: 10.3389/fragi.2024.1493406

Korea Association Resource (KARE) of the Korean Genome and Epidemiology Study (KoGES) Health Examinees (HEXA) of the KoGES

King-Hudson Te-Rina J., Andree G. Pearson<sup>1</sup>, Caitlin Dunstan-Harrison<sup>2</sup>, Mathew T. Powell<sup>2</sup>, Nicholas J. Magon<sup>1</sup>, Teagan S. Edwards<sup>1</sup>, Louise N. Paton<sup>1</sup>, Jeffry S. Tang<sup>1</sup>, Anthony J. Kettle<sup>1</sup>, John F. Pearson<sup>3</sup>, Jesse Kokaua<sup>4</sup>, Hayley Guiney<sup>4</sup>, Sandhya Ramrakha<sup>4</sup>, Richie Poulton<sup>4</sup>, Terrie E. Moffitt<sup>5,6</sup>, Elizabeth C. Ledgerwood<sup>1,2\*</sup> & Mark B. Hampton<sup>1\*</sup> (2025) Biomarkers of oxidative and mitochondrial stress are associated with accelerated pace of aging at midlife in a birth cohort.

Dunedin Study, in review

Knisely, M. R., Masese, R. V., Mathias, J. G., Yang, Q., Hatch, D., Lê, B. M., ... & Ashley-Koch, A. (2024). Epigenetic Aging Associations With Psychoneurological Symptoms and Social Functioning in Adults With Sickle Cell Disease. *Biological Research For Nursing*, 10998004241250322.

Kresovich, J. K., O'Brien, K. M., Xu, Z., Weinberg, C. R., Sandler, D. P., & Taylor, J. A. (2023). Abstract 5757: Breast cancer diagnosis and treatment associated with acceleration of biological aging over time in a racially diverse cohort of women. *Cancer Research*, 83(7\_Supplement), 5757-5757. <https://doi.org/10.1158/1538-7445.AM2023-5757>

Kresovich, J. K., O'Brien, K. M., Xu, Z., Weinberg, C. R., Sandler, D. P., & Taylor, J. A. (2023). Changes in methylation-based aging in women who do and do not develop breast cancer. *J Natl Cancer Inst*, 115(11), 1329-1336. <https://doi.org/10.1093/jnci/djad117>  
Sister Study

Kresovich, J. K., Sandler, D. P., & Taylor, J. A. (2023). Methylation-Based Biological Age and Hypertension Prevalence and Incidence. *Hypertension*, 80(6), 1213-1222. <https://doi.org/10.1161/hypertensionaha.122.20796>  
Sister Study

Kuo, P. L., Moore, A. Z., Lin, F. R., & Ferrucci, L. (2021). Epigenetic Age Acceleration and Hearing: Observations From the Baltimore Longitudinal Study of Aging. *Front Aging Neurosci*, 13, 790926. <https://doi.org/10.3389/fnagi.2021.790926>  
Baltimore Longitudinal Study of Aging

Lê, B. M., Hatch, D., Yang, Q., Shah, N., Luyster, F. S., Garrett, M. E., Tanabe, P., Ashley-Koch, A. E., & Knisely, M. R. (2024). Characterizing epigenetic aging in an adult sickle cell disease cohort. *Blood Adv*, 8(1), 47-55. <https://doi.org/10.1182/bloodadvances.2023011188>

Lee, Y., Bohlin, J., Page, C. M., Nustad, H. E., Harris, J. R., Magnus, P., Jugessur, A., Magnus, M. C., Häberg, S. E., & Hanevik, H. I. (2022). Associations between epigenetic age acceleration and infertility. *Hum Reprod*, 37(9), 2063-2074. <https://doi.org/10.1093/humrep/deac147>  
Norwegian MOBA

Lee, H. S., Kim, B., & Park, T. (2024). The association between sleep quality and accelerated epigenetic aging with metabolic syndrome in Korean adults. *Clinical Epigenetics*, 16(1), 92.

- Li, Marissa, Jenna A. Fernandez, Kristina Kirschner, Tamir Chandra, Kristen B. McCullough, Sarah Aug, Terra L. Lasho, Christy Finke, Naseema Gangat, Abhishek A. Mangaonkar, Rachel A Simon, Alejandro Ferrer, Mrinal M. Patnaik, James M. Foran, Cecilia Y. Arana Yi, Yael Kusne, Talha Badar (2024) A Prospective Cohort Study of Biologic and Chronologic Aging in 106 Patients with Age and Context-Relevant Clonal Hematopoiesis, *Blood*, Volume 144, Supplement 1, Page 5632, ISSN 0006-4971, <https://doi.org/10.1182/blood-2024-206069>.
- Li Piani, L., Vigano, P., & Somigliana, E. (2023). Epigenetic clocks and female fertility timeline: A new approach to an old issue? *Front Cell Dev Biol*, 11, 1121231. <https://doi.org/10.3389/fcell.2023.1121231>
- Lin, L., Kiryakos, J., Ammous, F., Ratliff, S. M., Ware, E. B., Faul, J. D., Kardia, S. L. R., Zhao, W., Birditt, K. S., & Smith, J. A. (2024). Epigenetic age acceleration is associated with blood lipid levels in a multi-ancestry sample of older U.S. adults. *Res Sq.* <https://doi.org/10.21203/rs.3.rs-3934965/v1>  
HRS
- Lin, W. Y. (2023). Epigenetic clocks derived from western samples differentially reflect Taiwanese health outcomes. *Front Genet*, 14, 1089819. <https://doi.org/10.3389/fgene.2023.1089819>  
Taiwanese BioBank
- Magnus, M.C., Lee, Y., Carlsen, E.Ø. et al. Parental epigenetic age acceleration and risk of adverse birth outcomes: the Norwegian mother, father and child cohort study. *BMC Med* 22, 554 (2024). <https://doi.org/10.1186/s12916-024-03780-7>  
Norwegian MOBA
- Mammadova, J., Palfi, S., Torriente, A. G., Adler, E., Cruz, R., AL-JUMAYLI, M., ... & Kresovich, J. (2024). PRETREATMENT METHYLATION-BASED BIOLOGICAL AGE IS A NOVEL RISK FACTOR FOR TRASTUZUMAB-INDUCED CARDIOTOXICITY. *Journal of the American College of Cardiology*, 83(13\_Supplement), 2324-2324.
- McMurran, C. E., Wang, Y., Mak, J. K. L., Karlsson, I. K., Tang, B., Ploner, A., Pedersen, N. L., & Hägg, S. (2023). Advanced biological ageing predicts future risk for neurological diagnoses and clinical examination findings. *Brain*, 146(12), 4891-4902. <https://doi.org/10.1093/brain/awad252>
- Meier, H. C. S., Mitchell, C., Karadimas, T., & Faul, J. D. (2023). Systemic inflammation and biological aging in the Health and Retirement Study. *Geroscience*, 45(6), 3257-3265. <https://doi.org/10.1007/s11357-023-00880-9>  
HRS
- Miao, K., Hong, X., Cao, W., Lv, J., Yu, C., Huang, T., Sun, D., Liao, C., Pang, Y., Hu, R., Pang, Z., Yu, M., Wang, H., Wu, X., Liu, Y., Gao, W., & Li, L. (2024). Association between epigenetic age and type 2 diabetes mellitus or glycemic traits: A longitudinal twin study. *Aging Cell*, n/a(n/a), e14175. <https://doi.org/https://doi.org/10.1111/acel.14175>  
Chinese National Twin Registry

Nannini Drew Robert et al., (2024) Chimeric antigen receptor T-cell therapy in B-cell malignancies and DNA-methylation-based biological aging.. *JCO* **42**, 2562-2562(2024). DOI:10.1200/JCO.2024.42.16\_suppl.2562

Nannini, D. R., Nandu, N., Hildebrandt, G., & Cortese, R. (2024). DNA methylation-based biological age and time to relapse in triple negative breast cancer.

Nicholson, T., Dhaliwal, A., Quinlan, J. I., Allen, S. L., Williams, F. R., Hazeldine, J., ... & Lord, J. M. (2024). Accelerated aging of skeletal muscle and the immune system in patients with chronic liver disease. *Experimental & Molecular Medicine*, **56**(7), 1667-1681.

Peterson, M. D., Collins, S., Meier, H. C. S., Brahmstedt, A., & Faul, J. D. (2023). Grip strength is inversely associated with DNA methylation age acceleration. *J Cachexia Sarcopenia Muscle*, **14**(1), 108-115. <https://doi.org/10.1002/jcsm.13110>  
HRS

Phyo, A. Z. Z., Fransquet, P. D., Wrigglesworth, J., Woods, R. L., Espinoza, S. E., & Ryan, J. (2024). Sex differences in biological aging and the association with clinical measures in older adults. *Geroscience*, **46**(2), 1775-1788. <https://doi.org/10.1007/s11357-023-00941-z>  
ASPREE

Poganik, J. R., Zhang, B., Baht, G. S., Tyshkovskiy, A., Deik, A., Kerepesi, C., Yim, S. H., Lu, A. T., Haghani, A., Gong, T., Hedman, A. M., Andolf, E., Pershagen, G., Almqvist, C., Clish, C. B., Horvath, S., White, J. P., & Gladyshev, V. N. (2023). Biological age is increased by stress and restored upon recovery. *Cell Metab*, **35**(5), 807-820.e805.  
<https://doi.org/10.1016/j.cmet.2023.03.015>

Three Clinical Datasets

Rasmussen, L. J. H., Caspi, A., Ambler, A., Danese, A., Elliott, M., Eugen-Olsen, J., Hariri, A. R., Harrington, H., Houts, R., Poulton, R., Ramrakha, S., Sugden, K., Williams, B., & Moffitt, T. E. (2021). Association Between Elevated suPAR, a New Biomarker of Inflammation, and Accelerated Aging. *J Gerontol A Biol Sci Med Sci*, **76**(2), 318-327.  
<https://doi.org/10.1093/gerona/glaa178>  
Dunedin Study

Rentscher, K. E., Bethea, T. N., Zhai, W., Small, B. J., Zhou, X., Ahles, T. A., Ahn, J., Breen, E. C., Cohen, H. J., Extermann, M., Graham, D. M. A., Jim, H. S. L., McDonald, B. C., Nakamura, Z. M., Patel, S. K., Root, J. C., Saykin, A. J., Van Dyk, K., Mandelblatt, J. S., & Carroll, J. E. (2023). Epigenetic aging in older breast cancer survivors and noncancer controls: preliminary findings from the Thinking and Living with Cancer Study. *Cancer*, **129**(17), 2741-2753. <https://doi.org/10.1002/cncr.34818>  
Thinking and Living with Cancer Study

Roach, K., Collins, C., Culp, E. V., Ogu, U. O., & Mozhui, K. (2024). Epigenetic Clocks and Accelerated Aging in Sickle Cell Disease. *Blood*, **144**, 3865.

Safaee, M. M., Dwaraka, V. B., Lee, J. M., Fury, M., Mendez, T. L., Smith, R., Lin, J., Smith, D. L., Burke, J. F., Scheer, J. K., Went, H., & Ames, C. P. (2024). Epigenetic age biomarkers and risk assessment in adult spinal deformity: a novel association of

biological age with frailty and disability. *J Neurosurg Spine*, 40(3), 312-323.  
<https://doi.org/10.3171/2023.10.Spine23435>

Sehgal, R., Meer, M., Shadyab, A. H., Casanova, R., Manson, J. E., Bhatti, P., Crimmins, E. M., Assimes, T. L., Whitsel, E. A., Higgins-Chen, A. T., & Levine, M. (2023). Systems Age: A single blood methylation test to quantify aging heterogeneity across 11 physiological systems. *bioRxiv*. <https://doi.org/10.1101/2023.07.13.548904>  
 Women's Health Initiative

Singh, S., Giron, L. B., Shaikh, M. W., Shankaran, S., Engen, P. A., Bogin, Z. R., Bambi, S. A., Goldman, A. R., Azevedo, J., Orgaz, L., de Pedro, N., González, P., Giera, M., Verhoeven, A., Sánchez-López, E., Pandrea, I. V., Kannan, T., Tanes, C. E., Bittinger, K., . . . Abdel-Mohsen, M. (2023). Distinct Intestinal Microbial Signatures Linked to Accelerated Biological Aging in People with HIV. *Res Sq.*  
<https://doi.org/10.21203/rs.3.rs-3492242/v1>

Sung, Hsien-Liang and Wan-Yu Lin. (2024). Causal effects of cardiovascular health on five epigenetic clocks, *Clinical Epigenetics* 16(1). 10.1186/s13148-024-01752-5  
 Taiwanese Biobank

Tamargo, J. A., Strath, L. J., & Cruz-Almeida, Y. (2024). High-impact pain is associated with epigenetic aging among middle-aged and older adults: Findings from the Health and Retirement Study. *The Journals of Gerontology, Series A: Biological Sciences and Medical Sciences*, glae149.  
 HRS

Tang, B., Li, X., Wang, Y., Sjölander, A., Johnell, K., Thambisetty, M., Ferrucci, L., Reynolds, C. A., Finkel, D., Jylhävä, J., Pedersen, N. L., & Hägg, S. (2023). Longitudinal associations between use of antihypertensive, antidiabetic, and lipid-lowering medications and biological aging. *Geroscience*, 45(3), 2065-2078. <https://doi.org/10.1007/s11357-023-00784-8>  
 Swedish Adoption/Twin Study of Aging (SATSA)

Torma, F., Kerepesi, C., Jókai, M., Babszki, G., Koltai, E., Ligeti, B., Kalcsevszki, R., McGreevy, K. M., Horvath, S., & Radák, Z. (2024). Alterations of the gut microbiome are associated with epigenetic age acceleration and physical fitness. *Aging Cell*, e14101.  
<https://doi.org/10.1111/acel.14101>

Verschoor, C. P., Vlasschaert, C., Rauh, M. J., & Paré, G. (2023). A DNA methylation based measure outperforms circulating CRP as a marker of chronic inflammation and partly reflects the monocytic response to long-term inflammatory exposure: A Canadian Longitudinal Study on Aging analysis. *Aging Cell*, 22(7), e13863.  
<https://doi.org/10.1111/acel.13863>  
 Canadian longitudinal study on aging (CLSA)

Vetter, V. M., Demircan, K., Homann, J., Chillon, T. S., Muelleder, M., Shomroni, O., . . . & Demuth, I. (2024). Low Blood Levels of Selenium, Selenoprotein P and GPx3 are Associated with Accelerated Biological Aging: Results from the Berlin Aging Study II (BASE-II). *medRxiv*, 2024-04.  
 Berlin Aging Study II (BASE-II).

Viglino, J., Casas-Recasen, S., Olvera, N., Garcia, T., Agusti, A., & Faner, R. (2024). Epigenetic age acceleration and severity of airflow limitation on blood and lung tissue of **COPD**. European Respiratory Journal.

Wang, H., Liu, Z., Fan, H., Guo, C., Zhang, X., Li, Y., ... & Zhang, T. (2024). Association between advanced fibrosis and epigenetic age acceleration among individuals with MASLD. *Journal of Gastroenterology*, 1-9.

Whitman E.T., Elliott, M.L., Knodt, A.R., Abraham, W.C., Anderson, T.J., Cutfield, N., Hogan, S., Ireland, D., Melzer, T.R., Ramrakha, S., Sugden, K., Theodore, R., Williams, B.S., Caspi, A., Moffitt, T.E., & Hariri, A.R. for the Alzheimer's Disease Neuroimaging Initiative. (Submitted 2024). An estimate of the longitudinal pace of aging from a single brain scan predicts dementia conversion, morbidity, and mortality.

ADNI, UKBiobank, BRAINLAT

Wikström Shemer, D., Mostafaei, S., Tang, B., Pedersen, N. L., Karlsson, I. K., Fall, T., & Hägg, S. (2024). Associations between epigenetic aging and diabetes mellitus in a Swedish longitudinal study. *GeroScience*, 46(5), 5003-5014.

Ying, Kejun, Seth Paulson, Alec Eames, Alexander Tyshkovskiy, Siyuan Li, Martin Perez-Guevara, Mehrnoosh Emamifar, Maximiliano Casas Martínez, Dayoon Kwon, Anna Kosheleva, Michael P. Snyder, Dane Gobel, Chiara Herzog, Jesse R. Paganik, Biomarker of Aging Consortium, Mahdi Moqri, Vadim N. Gladyshev. (2024 preprint) A Unified Framework for Systematic Curation and Evaluation of Aging Biomarkers.

<https://www.researchsquare.com/article/rs-4481437/v1.pdf>

Normative Aging Study, MassGeneral Brigham dataset

Ying, K., Tyshkovskiy, A., Chen, Q., Latorre-Crespo, E., Zhang, B., Liu, H., ... & Gladyshev, V. N. (2024). High-dimensional Ageome Representations of Biological Aging across Functional Modules. *bioRxiv*.

Normative Aging Study, MassGeneral Brigham Dataset

## PUBLICATIONS ON ALZHEIMERS, DEMENTIAS, COGNITIVE DECLINE, AND BRAIN OUTCOMES

Belsky, D. W., Caspi, A., Houts, R., Cohen, H. J., Corcoran, D. L., Danese, A., Harrington, H., Israel, S., Levine, M. E., Schaefer, J. D., Sugden, K., Williams, B., Yashin, A. I., Poulton, R., & Moffitt, T. E. (2015). Quantification of biological aging in young adults. *Proc Natl Acad Sci U S A*, 112(30), E4104-4110. <https://doi.org/10.1073/pnas.1506264112>  
Dunedin Study

Belsky, D. W., Moffitt, T. E., Cohen, A. A., Corcoran, D. L., Levine, M. E., Prinz, J. A., Schaefer, J., Sugden, K., Williams, B., Poulton, R., & Caspi, A. (2018). Eleven Telomere, Epigenetic Clock, and Biomarker-Composite Quantifications of Biological Aging: Do They Measure the Same Thing? *Am J Epidemiol*, 187(6), 1220-1230.  
<https://doi.org/10.1093/aje/kwx346>

Dunedin Study

Chen, J., Moubaddar, L., Clausing, E., Kezios, K., Conneely, K., Huls, A., Baccarelli, A., Factor-Litvak, P., Cirillo, P., Shelton, R., Link, B., & Suglia, S. (In Review). Associations of childhood, adolescence, and midlife cognitive function with DNA methylation age acceleration in midlife *Aging (Albany NY)*.

Child Health and Development Study

Elliott, M. L., Belsky, D. W., Knodt, A. R., Ireland, D., Melzer, T. R., Poulton, R., Ramrakha, S., Caspi, A., Moffitt, T. E., & Hariri, A. R. (2021). Brain-age in midlife is associated with accelerated biological aging and cognitive decline in a longitudinal birth cohort. *Mol Psychiatry*, 26(8), 3829-3838. <https://doi.org/10.1038/s41380-019-0626-7>

Dunedin Study

Elliott, M. L., Caspi, A., Houts, R. M., Ambler, A., Broadbent, J. M., Hancox, R. J., Harrington, H., Hogan, S., Keenan, R., Knodt, A., Leung, J. H., Melzer, T. R., Purdy, S. C., Ramrakha, S., Richmond-Rakerd, L. S., Righarts, A., Sugden, K., Thomson, W. M., Thorne, P. R., . . . Moffitt, T. E. (2021). Disparities in the pace of biological aging among midlife adults of the same chronological age have implications for future frailty risk and policy. *Nat Aging*, 1(3), 295-308. <https://doi.org/10.1038/s43587-021-00044-4>

Dunedin Study

Engvig, A., Kalleberg, K. T., Westlye, L. T., & Leonardsen, E. H. (2024). Complementary value of molecular, phenotypic, and functional aging biomarkers in dementia prediction. *GeroScience*, 1-20.

ADNI

Felt, J. M., Yusupov, N., Harrington, K. D., Fietz, J., Zhang, Z. Z., Sliwinski, M. J., Ram, N., O'Donnell, K. J., Meaney, M. J., Putnam, F. W., Noll, J. G., Binder, E. B., & Shenk, C. E. (2023). Epigenetic age acceleration as a biomarker for impaired cognitive abilities in adulthood following early life adversity and psychiatric disorders. *Neurobiol Stress*, 27, 100577. <https://doi.org/10.1016/j.ynstr.2023.100577>

BECOME Study, Female Growth and Development Study

Gonzales, HM. Et al. and Myriam Fornage. (AAIC poster 2024) Epigenetic Aging associations with plasma biomarkers.  
SOL-INCA, HCHS/SOL

Holmes, H. E., Valentin, R. E., Jernérén, F., de Jager Loots, C. A., Refsum, H., Smith, A. D., . . . & Alzheimer's Disease Neuroimaging Initiative. (2024). Elevated homocysteine is associated with increased rates of epigenetic aging in a population with mild cognitive impairment. *Aging Cell*, 23(10), e14255.

Karalija, N., Papenberg, G., Johansson, J., Wåhlin, A., Salami, A., Andersson, M., Axelsson, J., Kuznetsov, D., Riklund, K., Lövdén, M., Lindenberger, U., Bäckman, L., & Nyberg, L. (2024). Longitudinal support for the correlative triad among aging, dopamine D2-like receptor loss, and memory decline. *Neurobiol Aging*, 136, 125-132. <https://doi.org/10.1016/j.neurobiolaging.2024.02.001>  
Cognition, Brain, and Aging (COBRA)

Li, Zexu. Huitong Ding, Mengyao Wang, Yi Li, Ting Fang Alvin Ang, Gurnani Ashita, Gifford Katherine, Cody Karjadi, Daniel Levy, Rhoda Au, Chunyu Liu Association of DNA methylation age acceleration with digital clock drawing test performance: the Framingham Heart Study medRxiv 2024.11.06.24316862; doi: <https://doi.org/10.1101/2024.11.06.24316862>  
Framingham Heart Study

Nguyen, S., McEvoy, L. K., Espeland, M. A., Whitsel, E. A., Lu, A., Horvath, S., Manson, J. E., Rapp, S. R., & Shadyab, A. H. (2024). Associations of Epigenetic Age Estimators With Cognitive Function Trajectories in the Women's Health Initiative Memory Study. *Neurology*, 103(1), e209534. <https://doi.org/10.1212/WNL.0000000000209534>  
Women's Health Initiative Memory Study

Phyo, A. Z. Z., Wu, Z., Espinoza, S. E., Murray, A. M., Fransquet, P. D., Wrigglesworth, J., ... & Ryan, J. (2024). Epigenetic age acceleration and cognitive performance over time in older adults. *Alzheimer's & Dementia: Diagnosis, Assessment & Disease Monitoring*, 16(3), e70010.  
ASPREE

Piyush, G., Sai Pavan Kumar, V., Katja-Elisabeth, P., Reinhold, S., & Helena, S. (2023). Epigenetic age acceleration is related to cognition and cognitive decline in the Elderly: Results of the Austrian Stroke Prevention Study. *medRxiv*, 2023.2011.2021.23298753. <https://doi.org/10.1101/2023.11.21.23298753>  
Austrian Stroke Prevention Study

Reed, R. G., Carroll, J. E., Marsland, A. L., & Manuck, S. B. (2022). DNA methylation-based measures of biological aging and cognitive decline over 16-years: preliminary longitudinal findings in midlife. *Aging (Albany NY)*, 14(23), 9423-9444. <https://doi.org/10.1863/aging.204376>  
AHAB Study

Savin, M.J, Wang, H. Pei, H., Aiello, A.E., Assuras, S., Caspi, A., Moffitt, T.E., Muenning, P.A., Ryan, C.P., Shi, B. Stern, Y., Sugden, K., Valeri, L. Belsky, D.W. (2024). Faster DunedinPACE, an epigenetic clock for pace of biological aging, is associated with accelerated cognitive aging among older adults in the Framingham Heart Study. *Alzheimer's & Dementia: Diagnosis, Assessment and Disease Monitoring*.  
Framingham Heart Study

Schäfer Hackenhaar, F., Josefsson, M., Nordin Adolfsson, A., Landfors, M., Kauppi, K., Porter, T., Milicic, L., Laws, S. M., Hultdin, M., Adolfsson, R., Degerman, S., & Pudas, S. (2023). Sixteen-Year Longitudinal Evaluation of Blood-Based DNA Methylation Biomarkers for Early Prediction of Alzheimer's Disease. *J Alzheimers Dis*, 94(4), 1443-1464. <https://doi.org/10.3233/jad-230039>  
Australian Imaging Biomarkers and Lifestyle (AIBL), The Betula Study Sweden

Sugden, K., Caspi, A., Elliott, M. L., Bourassa, K. J., Chamarti, K., Corcoran, D. L., Hariri, A. R., Houts, R. M., Kothari, M., Kritchevsky, S., Kuchel, G. A., Mill, J. S., Williams, B. S., Belsky, D. W., & Moffitt, T. E. (2022). Association of Pace of Aging Measured by Blood-Based DNA Methylation With Age-Related Cognitive Impairment and Dementia. *Neurology*, 99(13), e1402-e1413. <https://doi.org/10.1212/WNL.0000000000200898>  
ADNI Study (Alzheimers Disease Neuroimaging Initiative), Framingham Offspring Study

Thomas, A., Ryan, C. P., Caspi, A., Moffitt, T. E., Sugden, K., Zhou, J., Belsky, D. W., & Gu, Y. (2023). Diet, pace of biological aging, and risk of dementia in the Framingham Heart Study. *medRxiv*. <https://doi.org/10.1101/2023.05.24.23290474>  
 Framingham Offspring Study

Whitman, E. T., Ryan, C. P., Abraham, W. C., Addae, A., Corcoran, D. L., Elliott, M. L., Hogan, S., Ireland, D., Keenan, R., Knodt, A. R., Melzer, T. R., Poulton, R., Ramrakha, S., Sugden, K., Williams, B. S., Zhou, J., Hariri, A. R., Belsky, D. W., Moffitt, T. E., & Caspi, A. (2023). A blood biomarker of accelerated aging in the body associates with worse structural integrity in the brain: replication across three cohorts. *medRxiv*.  
<https://doi.org/10.1101/2023.09.06.23295140>  
 ADNI, Framingham, Dunedin Study

Whitman E.T., Elliott, M.L., Knodt, A.R., Abraham, W.C., Anderson, T.J., Cutfield, N., Hogan, S., Ireland, D., Melzer, T.R., Ramrakha, S., Sugden, K., Theodore, R., Williams, B.S., Caspi, A., Moffitt, T.E., & Hariri, A.R. for the Alzheimer's Disease Neuroimaging Initiative. (Submitted 2024). An estimate of the longitudinal pace of aging from a single brain scan predicts dementia conversion, morbidity, and mortality.  
 ADNI, Dunedin Study, UK Biobank, BRAINLAT

Yannatos, I., Brown, R., Boen, C., Stites, S. D., Xie, S. X., & McMillan, C. T. (2023). Racial disparities in epigenetic and cognitive aging. *Alzheimer's & Dementia*, 19(S22), e078504. <https://doi.org/https://doi.org/10.1002/alz.078504>  
 HRS

Yannatos, I., Stites, S. D., Boen, C., Xie, S. X., Brown, R. T., & McMillan, C. T. (2023). Epigenetic age and socioeconomic status contribute to racial disparities in cognitive and functional aging between Black and White older Americans. *medRxiv*.  
<https://doi.org/10.1101/2023.09.29.23296351>  
 HRS

Ying, Kejun, Seth Paulson, Alec Eames, Alexander Tyshkovskiy, Siyuan Li, Martin Perez-Guevara, Mehrnoosh Emamifar, Maximiliano Casas Martínez, Dayoon Kwon, Anna Kosheleva, Michael P. Snyder, Dane Gobel, Chiara Herzog, Jesse R. Paganik, Biomarker of Aging Consortium, Mahdi Moqri, Vadim N. Gladyshev. (2024 preprint) A Unified Framework for Systematic Curation and Evaluation of Aging Biomarkers.

<https://www.researchsquare.com/article/rs-4481437/v1.pdf>

Normative Aging Study, MassGeneral Brigham dataset

#### PUBLICATIONS ON Health Inequalities:

Aikins, M., Willems, Y., C, M., Goosby, B., & Raffington, L. (2023). Associations of racialized inequities with saliva DNA-methylation measures of biological aging and mental health across childhood and adolescence. <https://doi.org/10.17605/OSF.IO/V49KC>  
 Future of Families and Child Wellbeing Study

- Andrasfay, T., & Crimmins, E. (2023). Occupational characteristics and epigenetic aging among older adults in the United States. *Epigenetics*, 18(1), 2218763.  
<https://doi.org/10.1080/15592294.2023.2218763>  
HRS
- Avila-Rieger, J., Turney, I. C., Vonk, J. M. J., Esie, P., Seblova, D., Weir, V. R., Belsky, D. W., & Manly, J. J. (2022). Socioeconomic Status, Biological Aging, and Memory in a Diverse National Sample of Older US Men and Women. *Neurology*, 99(19), e2114-e2124.  
<https://doi.org/10.1212/wnl.00000000000201032>  
HRS
- Bao, Y., Gorrie-Stone, T., Hannon, E., Hughes, A., Andrayas, A., Neilson, G., Burrage, J., Mill, J., Schalkwyk, L., & Kumari, M. (2022). Social mobility across the lifecourse and DNA methylation age acceleration in adults in the UK. *Sci Rep*, 12(1), 22284.  
<https://doi.org/10.1038/s41598-022-26433-2>  
HRS
- Bourassa, K. J., Halverson, T. F., Garrett, M. E., Hair, L., Dennis, M., Ashley-Koch, A. E., Beckham, J. C., & Kimbrel, N. A. (2024). Demographic characteristics and epigenetic biological aging among post-9/11 veterans: Associations of DunedinPACE with sex, race, and age. *Psychiatry Res*, 336, 115908.  
<https://doi.org/10.1016/j.psychres.2024.115908>  
PDMH Veteran's Study
- Berg, M. T., Riley, K., Lei, M. K., & Simons, R. L. (2024). Incarceration Exposure, Biological Aging, and Depression Symptoms in an African American Sample of Older Adults. *Journal of aging and health*, 08982643241257065.  
FACHS
- Chang OD, Meier HCS, Maguire-Jack K, Davis-Kean P, Mitchell C. Childhood Maltreatment and Longitudinal Epigenetic Aging: NIMHD Social Epigenomics Program. *JAMA Netw Open*. 2024;7(7):e2421877. doi:10.1001/jamanetworkopen.2024.21877  
Future of Families and Child Wellbeing
- Clair, A., Baker, E., & Kumari, M. (2023). Are housing circumstances associated with faster epigenetic ageing? *J Epidemiol Community Health*, 78(1), 40-46.  
<https://doi.org/10.1136/jech-2023-220523>  
UK Household Longitudinal Study
- Crimmins, E. M., Thyagarajan, B., Levine, M. E., Weir, D. R., & Faul, J. (2021). Associations of Age, Sex, Race/Ethnicity, and Education With 13 Epigenetic Clocks in a Nationally Representative U.S. Sample: The Health and Retirement Study. *J Gerontol A Biol Sci Med Sci*, 76(6), 1117-1123. <https://doi.org/10.1093/gerona/glab016>  
HRS
- Choi, Eun Young and Jennifer A Ailshire, Neighborhood Stressors and Epigenetic Age Acceleration Among Older Americans, *The Journals of Gerontology: Series B*, Volume 79, Issue 12, December 2024, gbae176, <https://doi.org/10.1093/geronb/gbae176>  
HRS

**Crimmins:** Crimmins, E. M., Klopack, E. T., & Kim, J. K. (2024). Generations of epigenetic clocks and their links to **socioeconomic status** in the Health and Retirement Study. *Epigenomics*, 1–12. <https://doi.org/10.1080/17501911.2024.2373682>  
The Health and Retirement Study (HRS)

Cuevas, A. G., Cole, S. W., Belsky, D. W., McSorley, A.-M., Shon, J. M., & Chang, V. W. (2024). Multi-discrimination exposure and biological aging: Results from the midlife in the United States study. *Brain, Behavior, & Immunity - Health*, 100774. <https://doi.org/https://doi.org/10.1016/j.bbih.2024.100774>  
MIDUS

Das, A. (2023). Retirement and Epigenetic age Acceleration Among Older U.S. Adults. *Adaptive Human Behavior and Physiology*, 9(3), 264-283. <https://doi.org/10.1007/s40750-023-00221-2>  
HRS

Del Toro, J., Freilich, C., Rea-Sandin, G., Markon, K., Wilson, S., & Krueger, R. (2023). Ethnic/racial disparities in DNA methylation age profiles across the lifespan. Future of Families and Child Wellbeing (FFCW) study, Midlife in the United States Study (MIDUS)

Del Toro J, Martz C, Freilich CD, et al. (2024) Longitudinal Changes in Epigenetic Age Acceleration Across Childhood and Adolescence. *JAMA Pediatr.* 2024;178(12):1298–1306. doi:10.1001/jamapediatrics.2024.3669  
Future of Families and Child Wellbeing (FFCW) study,

Engelman, M. (2023). *Neighborhood, race, and epigenetic ageing clocks*. Duke University, Durham, NC.  
REWARD Study, Wisconsin

Freni-Sterrantino, A., Fiorito, G., D'Errico, A., Robinson, O., Virtanen, M., Ala-Mursula, L., Järvelin, M. R., Ronkainen, J., & Vineis, P. (2022). Work-related stress and well-being in association with epigenetic age acceleration: A Northern Finland Birth Cohort 1966 Study. *Aging (Albany NY)*, 14(3), 1128-1156. <https://doi.org/10.18632/aging.203872>  
Northern Finland Birth Cohort 1966 Study

Freni-Sterrantino, A., Fiorito, G., d'Errico, A., Virtanen, M., Ala-Mursula, L., Järvelin, M. R., Vineis, P., & Robinson, O. (2022). Association between work characteristics and epigenetic age acceleration: cross-sectional results from UK - Understanding Society study. *Aging (Albany NY)*, 14(19), 7752-7773. <https://doi.org/10.18632/aging.204327>  
Understanding Society Study

Graf, G. H., Crowe, C. L., Kothari, M., Kwon, D., Manly, J. J., Turney, I. C., Valeri, L., & Belsky, D. W. (2022). Testing Black-White Disparities in Biological Aging Among Older Adults in the United States: Analysis of DNA-Methylation and Blood-Chemistry Methods. *Am J Epidemiol.* 191(4), 613-625. <https://doi.org/10.1093/aje/kwab281>  
HRS

- Graf, G. H., Zhang, Y., Domingue, B. W., Harris, K. M., Kothari, M., Kwon, D., Muennig, P., & Belsky, D. W. (2022) Social mobility and biological aging among older adults in the United States, 81(2), pgac029. <https://doi.org/10.1093/pnasnexus/pgac029>  
HRS
- Graf, G. H. J., Aiello, A. E., Caspi, A., Kothari, M., Liu, H., Moffitt, T. E., Muennig, P. A., Ryan, C. P., Sugden, K., & Belsky, D. W. (2024). Educational Mobility, Pace of Aging, and Lifespan Among Participants in the Framingham Heart Study. *JAMA Netw Open*, 7(3), e240655. <https://doi.org/10.1001/jamanetworkopen.2024.0655>  
The Framingham Offspring Study
- Harris, Kathleen Mullan, Brandt, Levitt., Gaydosh, Lauren, G., Chantel, M., Jess, M. M., Aura Ankita, M., Audrey, L. K., & Allison, E. A. (2024). Sociodemographic and Lifestyle Factors and Epigenetic Aging in US Young Adults: NIMHD Social Epigenomics Program. *JAMA Netw Open*. 2024;7(7):e2427889.  
doi:10.1001/jamanetworkopen.2024.27889  
Add Health
- Holuka, C., Menta, G., Caro, J. C., Vögele, C., D'Ambrosio, C., & Turner, J. D. (2024). Developmental epigenomic effects of maternal financial problems. *Development and Psychopathology*, 1-14.  
ALSPAC
- Joyce, B. T., Gao, T., Koss, K., Zheng, Y., Cardenas, A., Heiss, J., Just, A., Zhang, K., van Horn, L., Allen, N. B., Greenland, P., Cohen, S., Gordon-Larsen, P., Mitchell, C., McLanahan, S., Schneper, L., Notterman, D., Rifas-Shiman, S. L., Oken, E., . . . Hou, L. (2022). Impact of paternal education on epigenetic ageing in adolescence and mid-adulthood: a multi-cohort study in the USA and Mexico. *Int J Epidemiol*, 51(3), 870-884. <https://doi.org/10.1093/ije/dyab196>  
CARDIA, Fragile Families, Project Viva, Progress
- Kim, K., Joyce, B. T., Nannini, D. R., Zheng, Y., Gordon-Larsen, P., Shikany, J. M., Lloyd-Jones, D. M., Hu, M., Nieuwenhuijsen, M. J., Vaughan, D. E., Zhang, K., & Hou, L. (2023). Inequalities in urban greenness and epigenetic aging: Different associations by race and neighborhood socioeconomic status. *Sci Adv*, 9(26), eadf8140. <https://doi.org/10.1126/sciadv.adf8140>  
Coronary Artery Risk Development in Young Adults (CARDIA)
- Korous, K. M., Surachman, A., Rogers, C. R., & Cuevas, A. G. (2023). Parental education and epigenetic aging in middle-aged and older adults in the United States: A life course perspective. *Soc Sci Med*, 333, 116173. <https://doi.org/10.1016/j.socscimed.2023.116173>  
HRS
- Krieger N, Testa C, Chen JT, Johnson, N., Watkins, S. H., Suderman, M., Simpkin, A. J., Tilling, K., Waterman, P. D., Coull, B. A., De Vivo, I., Smith, G. D., Roux, A. V. D., & Relton, C. Epigenetic Aging and Racialized, Economic, and Environmental Injustice: NIMHD Social Epigenomics Program. *JAMA Netw Open*. 2024;7(7):e2421832.  
doi:10.1001/jamanetworkopen.2024.21832  
US My Body My Story, MESA Multi-Ethnic Atherosclerosis Study

Lei, M. K., & Beach, S. R. H. (2023). Neighborhood disadvantage is associated with biological aging: Intervention-induced enhancement of couple functioning confers resilience. *Fam Process*, 62(2), 818-834. <https://doi.org/10.1111/famp.12808>  
 The FACHS Study

Lei, M. K., Berg, M. T., Simons, R. L., & Beach, S. R. H. (2022). Neighborhood structural disadvantage and biological aging in a sample of Black middle age and young adults. *Soc Sci Med*, 293, 114654. <https://doi.org/10.1016/j.socscimed.2021.114654>  
 The FACHS Study

Lei, M. K., Lavner, J. A., Carter, S. E., Adesogan, O., & Beach, S. R. H. (2022). Relationship intervention indirectly buffers financial strain's effect on biological aging among Black adults. *J Fam Psychol*, 36(4), 502-512. <https://doi.org/10.1037/fam0000926>  
 The FACHS Study

Markon, K. E., Mann, F., Freilich, C., Cole, S., & Krueger, R. F. (2024). Associations between epigenetic age acceleration and longitudinal measures of psychosocioeconomic stress and status. *Social Science & Medicine*, 352, 116990.  
<https://doi.org/https://doi.org/10.1016/j.socscimed.2024.116990>  
 MIDUS

Martz, C. D., Benner, A. D., Goosby, B. J., Mitchell, C., & Gaydosh, L. (2024). Structural racism in primary schools and changes in epigenetic age acceleration among Black and White youth. *Social Science & Medicine*, 347, 116724.  
<https://doi.org/https://doi.org/10.1016/j.socscimed.2024.116724>  
 Future of Families and Child Wellbeing (FFCW) study,

Maunakea, A. K., Phankitnirundorn, K., Peres, R., Dye, C., Juarez, R., Walsh, C., ... & Le Marchand, L. (2024). Socioeconomic status, lifestyle, and DNA methylation age among racially and ethnically diverse adults: NIMHD Social Epigenomics Program. *JAMA Network Open*, 7(7), e2421889-e2421889.

Petrovic, D., Carmeli, C., Sandoval, J. L., Bodinier, B., Chadeau-Hyam, M., Schrempf, S., Ehret, G., Dhayat, N. A., Ponte, B., Pruijm, M., Vineis, P., Gonseth-Nusslé, S., Guessous, I., McCrory, C., Bochud, M., & Stringhini, S. (2023). Life-course socioeconomic factors are associated with markers of epigenetic aging in a population-based study. *Psychoneuroendocrinology*, 147, 105976.  
<https://doi.org/10.1016/j.psyneuen.2022.105976>  
 Swiss Family Study

Plascak, J. J., Archer, K., Llanos, A. A., Mooney, S. J., Xing, C. Y., Rundle, A. G., ... & Hong, C. C. (2024). Association between neighborhood disinvestment and all-cause survival moderated by epigenetic age among women with breast cancer. *Cancer Research*, 84(6\_Supplement), 6128-6128.

Raffington, L., Schwaba, T., Aikins, M., Richter, D., Wagner, G. G., Harden, K. P., Belsky, D. W., & Tucker-Drob, E. M. (2023). Associations of socioeconomic disparities with buccal DNA-methylation measures of biological aging. *Clin Epigenetics*, 15(1), 70.  
<https://doi.org/10.1186/s13148-023-01489-7>  
 SOEP-G Germany

Raffington, L., Tanksley, P. T., Sabhlok, A., Vinnik, L., Mallard, T., King, L. S., Goosby, B., Harden, K. P., & Tucker-Drob, E. M. (2023). Socially Stratified Epigenetic Profiles Are Associated With Cognitive Functioning in Children and Adolescents. *Psychol Sci*, 34(2), 170-185. <https://doi.org/10.1177/09567976221122760>  
Texas Twin Study

Raffington, L., Tanksley, P. T., Vinnik, L., Sabhlok, A., Patterson, M. W., Mallard, T., Malanchini, M., Ayorech, Z., Tucker-Drob, E. M., & Paige Harden, K. (2023). Associations of DNA-Methylation Measures of Biological Aging With Social Disparities in Child and Adolescent Mental Health. *Clinical Psychological Science*, 21677026231186802. <https://doi.org/10.1177/21677026231186802>  
Texas Twin Study

Schmitz, L. L., Zhao, W., Ratliff, S. M., Goodwin, J., Miao, J., Lu, Q., Guo, X., Taylor, K. D., Ding, J., Liu, Y., Levine, M., & Smith, J. A. (2022). The Socioeconomic Gradient in Epigenetic Ageing Clocks: Evidence from the Multi-Ethnic Study of Atherosclerosis and the Health and Retirement Study. *Epigenetics*, 17(6), 589-611. <https://doi.org/10.1080/15592294.2021.1939479>  
MESA, HRS

Shen, B., Mode, N. A., Noren Hooten, N., Pacheco, N. L., Ezike, N., Zonderman, A. B., & Evans, M. K. (2023). Association of Race and Poverty Status With DNA Methylation-Based Age. *JAMA Netw Open*, 6(4), e236340. <https://doi.org/10.1001/jamanetworkopen.2023.6340>  
The HANDLE Study

Simons, R. L., Lei, M.-K., Klopach, E., Berg, M., Zhang, Y., & Beach, S. S. R. (2021). (Re)Setting Epigenetic Clocks: An Important Avenue Whereby Social Conditions Become Biologically Embedded across the Life Course. *Journal of Health and Social Behavior*, 62(3), 436-453. <https://doi.org/10.1177/00221465211009309>  
The FACHS Study

Simons, R. L., Ong, M. L., Lei, M. K., Klopach, E., Berg, M., Zhang, Y., Philibert, R., Gibbons, F. X., & Beach, S. R. H. (2022). Shifts in lifestyle and socioeconomic circumstances predict change-for better or worse-in speed of epigenetic aging: A study of middle-aged black women. *Soc Sci Med*, 307, 115175. <https://doi.org/10.1016/j.soscimed.2022.115175>  
The FACHS Study

Smith AK, Katrinli S, Cobb DO, et al. Epigenetic Age Acceleration and Disparities in Posttraumatic Stress in Women in Southeast Louisiana: NIMHD Social Epigenomics Program. *JAMA Netw Open*. 2024;7(7):e2421884. doi:10.1001/jamanetworkopen.2024.21884  
Women and their Children's Health Cohort, Southeast Louisiana

Sugden, K., Moffitt, T. E., Arpwong, T. E., Arseneault, L., Belsky, D. W., Corcoran, D. L., Crimmins, E. M., Hannon, E., Houts, R., Mill, J. S., Poulton, R., Ramrakha, S., Wertz, J., Williams, B. S., & Caspi, A. (2023). Cross-National and Cross-Generational Evidence That Educational Attainment May Slow the Pace of Aging in European-Descent Individuals. *J Gerontol B Psychol Sci Soc Sci*, 78(8), 1375-1385. <https://doi.org/10.1093/geronb/gbad056>

Health and Retirement Study, Generation Scotland, E-Risk Study, UK Understanding Society, Dunedin Study

Surachman, A., Hamlat, E., Zannas, A. S., Horvath, S., Laraia, B., & Epel, E. (2024). Grandparents' educational attainment is associated with grandchildren's epigenetic-based age acceleration in the National Growth and Health Study. *Social Science & Medicine*, 355, 117142.

National Growth and Health Study

Tamargo, J. A., & Cruz-Almeida, Y. (2024). Food insecurity and epigenetic aging in middle-aged and older adults. *Social Science & Medicine*, 350, 116949.

HRS

Wang, W., Bao, Y., Haworth, S., & Kumari, M. (2024). Cross-sectional association between area deprivation and biological ageing: evidence from the UK Household Longitudinal Study. *The Lancet*, 404, S41.

UK Household Longitudinal Study

Ware, E. B., Higgins Tejera, C., Wang, H., Harris, S., Fisher, J. D., & Bakulski, K. M. (2024). Interplay of education and DNA methylation age on cognitive impairment: insights from the Health and Retirement Study. *GeroScience*, 1-14.

Yannatos, I., Brown, R., Boen, C., Stites, S. D., Xie, S. X., & McMillan, C. T. (2023). Racial disparities in epigenetic and cognitive aging. *Alzheimer's & Dementia*, 19(S22), e078504. <https://doi.org/https://doi.org/10.1002/alz.078504>

HRS

Yannatos, I., Stites, S. D., Boen, C., Xie, S. X., Brown, R. T., & McMillan, C. T. (2023). Epigenetic age and socioeconomic status contribute to racial disparities in cognitive and functional aging between Black and White older Americans. *medRxiv*. <https://doi.org/10.1101/2023.09.29.23296351>

HRS

Yusipov, Igor, Alena Kalyakulina, Claudio Franceschi, Mikhail Ivanchenko,  
**Map of epigenetic age acceleration:** A worldwide meta-analysis .

<https://www.biorxiv.org/content/biorxiv/early/2024/03/17/2024.03.17.585398.full.pdf>

DNAmethylation data from 93 datasets, 23K individuals' samples, 25 countries, and 31 ethnicities around the world

### PUBLICATIONS ON EARLY-LIFE RISK FACTORS:

Aiello, A. E., Mishra, A. A., Martin, C. L., Levitt, B., Gaydosh, L., Belsky, D. W., ... & Harris, K. M. (2024). Familial loss of a loved one and biological aging: NIMHD Social Epigenomics Program. *JAMA Network Open*, 7(7), e2421869-e2421869.

National Longitudinal Study of Adolescent to Adult Health

Belsky, D. W., Caspi, A., Cohen, H. J., Kraus, W. E., Ramrakha, S., Poulton, R., & Moffitt, T. E. (2017). Impact of early personal-history characteristics on the Pace of Aging:

implications for clinical trials of therapies to slow aging and extend healthspan. *Aging Cell*, 16(4), 644-651. <https://doi.org/10.1111/acel.12591>  
 Dunedin Study

Belsky, D. W., Caspi, A., Arseneault, L., Baccarelli, A., Corcoran, D. L., Gao, X., Hannon, E., Harrington, H. L., Rasmussen, L. J., Houts, R., Huffman, K., Kraus, W. E., Kwon, D., Mill, J., Pieper, C. F., Prinz, J. A., Poulton, R., Schwartz, J., Sugden, K., . . . Moffitt, T. E. (2020). Quantification of the pace of biological aging in humans through a blood test, the DunedinPoAm DNA methylation algorithm. *eLife*, 9. <https://doi.org/10.7554/eLife.54870>  
 E-Risk Study

Belsky, D. W., Caspi, A., Corcoran, D. L., Sugden, K., Poulton, R., Arseneault, L., Baccarelli, A., Chamarti, K., Gao, X., Hannon, E., Harrington, H. L., Houts, R., Kothari, M., Kwon, D., Mill, J., Schwartz, J., Vokonas, P., Wang, C., Williams, B. S., & Moffitt, T. E. (2022). DunedinPACE, a DNA methylation biomarker of the pace of aging. *eLife*, 11. <https://doi.org/10.7554/eLife.73420>  
 E-Risk Study

Bourassa, K. J., Moffitt, T. E., Ambler, A., Hariri, A. R., Harrington, H., Houts, R. M., Ireland, D., Knott, A., Poulton, R., Ramrakha, S., & Caspi, A. (2022). Association of Treatable Health Conditions During Adolescence With Accelerated Aging at Midlife. *JAMA Pediatr*, 176(4), 392-399. <https://doi.org/10.1001/jamapediatrics.2021.6417>  
 Dunedin Study

Brown RL, Alegria KE, Hamlat E, et al. Psychosocial Disadvantage During Childhood and Midlife Health: NIMHD Social Epigenomics Program. *JAMA Netw Open*. 2024;7(7):e2421841. doi:10.1001/jamanetworkopen.2024.21841  
 National Heart, Lung, and Blood Institute Growth and Health Study

Chang OD, Meier HCS, Maguire-Jack K, Davis-Kean P, Mitchell C. Childhood Maltreatment and Longitudinal Epigenetic Aging: NIMHD Social Epigenomics Program. *JAMA Netw Open*. 2024;7(7):e2421877. doi:10.1001/jamanetworkopen.2024.21877  
 Future of Families and Child Wellbeing Study

Cheng, M., Conley, D., Kuipers, T., Li, C., Ryan, C., Taubert, J., Wang, S., Wang, T., Zhou, J., Schmitz, L. L., Tobi, E. W., Heijmans, B., Lumey, L. H., & Belsky, D. W. (2023). Accelerated biological aging six decades after prenatal famine exposure. *medRxiv*. <https://doi.org/10.1101/2023.11.03.23298046>  
 Dutch Hunger Winter Study

DiMarzio, Karissa , Darlynn M. Rojo-Wissar, Evelyn Hernandez Valencia, Mikayla Ver Pault, Shane Denherder, Adamari Lopez, Jena Lerch, Georgette Metrailler, Sarah M. Merrill, April Highlander, Justin Parent Childhood Adversity and Adolescent Epigenetic Age Acceleration: The Role of Adolescent Sleep Health  
 medRxiv 2024.09.02.24312939; doi: <https://doi.org/10.1101/2024.09.02.24312939>  
 Future of Families and Child Wellbeing Study

Dye, C. K., Alschuler, D. M., Wu, H., Duarte, C., Monk, C., Belsky, D. W., ... & Scorza, P. (2024). Maternal adverse childhood experiences and biological aging during pregnancy and in newborns. *JAMA network open*, 7(8), e2427063-e2427063

- Farina, M. P., Klopack, E., Umberson, D., & Crimmins, E. (2024). The embodiment of parental death in early life through accelerated epigenetic aging: Implications for understanding how parental death before 18 shapes age-related health risk among older adults. *SSM - Population Health*, 101648. <https://doi.org/https://doi.org/10.1016/j.ssmph.2024.101648>  
HRS
- Felt, J. M., Harrington, K. D., Ram, N., O'Donnell, K. J., Sliwinski, M. J., Benson, L., Zhang, Z., Meaney, M. J., Putnam, F. W., Noll, J. G., & Shenk, C. E. (2023). Receptive Language Abilities for Females Exposed to Early Life Adversity: Modification by Epigenetic Age Acceleration at Midlife in a 30-Year Prospective Cohort Study. *J Gerontol B Psychol Sci Soc Sci*, 78(4), 585-595. <https://doi.org/10.1093/geronb/gbac158>  
Female Growth and Development Study
- Felt, J. M., Yusupov, N., Harrington, K. D., Fietz, J., Zhang, Z. Z., Sliwinski, M. J., Ram, N., O'Donnell, K. J., Meaney, M. J., Putnam, F. W., Noll, J. G., Binder, E. B., & Shenk, C. E. (2023). Epigenetic age acceleration as a biomarker for impaired cognitive abilities in adulthood following early life adversity and psychiatric disorders. *Neurobiol Stress*, 27, 100577. <https://doi.org/10.1016/j.ynstr.2023.100577>  
BECOME Study, Female Growth and Development Study
- Gonggrijp, B. M. A., Weijer, S. G. A. v. d., Bijleveld, C. C. J. H., Boomsma, D. I., & Dongen, J. v. (2024). Negative Life Events and Epigenetic Ageing: a Study in the Netherlands Twin Register. *bioRxiv*, 2024.2002.2020.581138. <https://doi.org/10.1101/2024.02.20.581138>  
Netherlands Twin Registry
- Hamlat, E. J., Moffitt, T. E., Meyer, A., Surachman, A., Dutcher, E., Laraia, B., Zhang, J., Lu, A. T., Horvath, S., & Epel, E. S. (In Review). Childhood Adversity of Mother Accelerates Offspring Biological Aging. *National Heart, Lung, and Blood Institute Growth and Health Study (NGHS, 1992)*.  
National Heart, Lung, and Blood Institute Growth and Health Study
- Hamlat, E. J., Neilands, T. B., Laraia, B., Zhang, J., Lu, A. T., Lin, J., Horvath, S., & Epel, E. S. (2023). Early life adversity predicts an accelerated cellular aging phenotype through early timing of puberty. *Psychol Med*, 53(16), 7720-7728.  
<https://doi.org/10.1017/s0033291723001629>  
National Heart, Lung, and Blood Institute Growth and Health Study
- Joshi, D., Gonzalez, A., Lin, D., & Raina, P. (2023). The association between adverse childhood experiences and epigenetic age acceleration in the Canadian longitudinal study on aging (CLSA). *Aging Cell*, 22(2), e13779. <https://doi.org/10.1111/acel.13779>  
Canadian Longitudinal Study on Aging
- Kim, J. K., Arpwong, T. E., Klopack, E. T., & Crimmins, E. M. (2023). Parental Divorce in Childhood and the Accelerated Epigenetic Aging for Earlier and Later Cohorts: Role of Mediators of Chronic Depressive Symptoms, Education, Smoking, Obesity, and Own Marital Disruption. *Journal of Population Ageing*. <https://doi.org/10.1007/s12062-023-09434-5>  
HRS

- Kim, K., Yaffe, K., Rehkopf, D. H., Zheng, Y., Nannini, D. R., Perak, A. M., Nagata, J. M., Miller, G. E., Zhang, K., Lloyd-Jones, D. M., Joyce, B. T., & Hou, L. (2023). Association of Adverse Childhood Experiences With Accelerated Epigenetic Aging in Midlife. *JAMA Netw Open*, 6(6), e2317987. <https://doi.org/10.1001/jamanetworkopen.2023.17987>  
CARDIA
- Klopack, E. T., Crimmins, E. M., Cole, S. W., Seeman, T. E., & Carroll, J. E. (2022). Accelerated epigenetic aging mediates link between adverse childhood experiences and depressive symptoms in older adults: Results from the Health and Retirement Study. *SSM Popul Health*, 17, 101071. <https://doi.org/10.1016/j.ssmph.2022.101071>  
HRS
- Kuzawa, C. W., Ryan, C. P., Adair, L. S., Lee, N. R., Carba, D. B., MacIsaac, J. L., Dever, K., Atashzay, P., Kobor, M. S., & McDade, T. W. (2022). Birth weight and maternal energy status during pregnancy as predictors of epigenetic age acceleration in young adults from metropolitan Cebu, Philippines. *Epigenetics*, 17(11), 1535-1545.  
<https://doi.org/10.1080/15592294.2022.2070105>  
Cebu Longitudinal Health and Nutrition Survey (CLHNS Phillipines)
- McCrory, C., Fiorito, G., O'Halloran, A. M., Polidoro, S., Vineis, P., & Kenny, R. A. (2022). Early life adversity and age acceleration at mid-life and older ages indexed using the next-generation GrimAge and Pace of Aging epigenetic clocks. *Psychoneuroendocrinology*, 137, 105643. <https://doi.org/10.1016/j.psyneuen.2021.105643>  
Swiss Family Study
- McDade, T. W., Ryan, C. P., Adair, L. S., Lee, N. R., Carba, D. B., MacIsaac, J. L., Dever, K., Atashzay, P., Kobor, M. S., & Kuzawa, C. W. (2023). Association between infectious exposures in infancy and epigenetic age acceleration in young adulthood in metropolitan Cebu, Philippines. *Am J Hum Biol*, 35(11), e23948. <https://doi.org/10.1002/ajhb.23948>  
CEBU
- Metrailler, G., Tavares, K., Ver Pault, M., Lopez, A., Denherder, S., Hernandez Valencia, E., ... & Parent, J. (2024). Community Threat, Positive Parenting, and Accelerated Epigenetic Aging: Longitudinal Links from Childhood to Adolescence. *medRxiv*, 2024-12.  
Future of Families and Child Wellbeing Study
- Niccodemi, G., Menta, G., Turner, J., & D'Ambrosio, C. (2022). Pace of aging, family environment and cognitive skills in children and adolescents. *SSM Popul Health*, 20, 101280. <https://doi.org/10.1016/j.ssmph.2022.101280>  
ALSPAC
- Perret, L. C., Geoffroy, M. C., Barr, E., Parnet, F., Provencal, N., Boivin, M., O'Donnell, K. J., Suderman, M., Power, C., Turecki, G., & Ouellet-Morin, I. (2022). Associations between epigenetic aging and childhood peer victimization, depression, and suicidal ideation in adolescence and adulthood: A study of two population-based samples. *Front Cell Dev Biol*, 10, 1051556. <https://doi.org/10.3389/fcell.2022.1051556>  
Quebec Longitudinal Study of Child Development, 1958 British Birth cohort

Qiao, X., Straight, B., Ngo, D., Hilton, C. E., Owuor Olungah, C., Naugle, A., ... & Needham, B. L. (2024). Severe drought exposure in utero associates to children's epigenetic age acceleration in a global climate change hot spot. *Nature Communications*, 15(1), 4140.

Richmond-Rakerd, L. S., Caspi, A., Ambler, A., d'Arbeloff, T., de Bruine, M., Elliott, M., Harrington, H., Hogan, S., Houts, R. M., Ireland, D., Keenan, R., Knodt, A. R., Melzer, T. R., Park, S., Poulton, R., Ramrakha, S., Rasmussen, L. J. H., Sack, E., Schmidt, A. T., . . . Moffitt, T. E. (2021). Childhood self-control forecasts the pace of midlife aging and preparedness for old age. *Proc Natl Acad Sci U S A*, 118(3).  
<https://doi.org/10.1073/pnas.2010211118>  
Dunedin Study

Ruiz, B., Broadbent, J. M., Thomson, W. M., Ramrakha, S., Moffitt, T. E., Caspi, A., & Poulton, R. (2023). Childhood caries is associated with poor health and a faster pace of aging by midlife. *J Public Health Dent*, 83(4), 381-388. <https://doi.org/10.1111/jphd.12591>  
Dunedin Study

Aino Saarinen, Aino, Saara M, Mishra Pashupati P, Leo-Pekka L, Binisha HM, Emma R, Nina M, Mika K, Olli R, Terho L, Liisa KJ. Early resilience and epigenetic ageing: Results from the prospective Young Finns Study with a 31-year follow-up. *Aging Cell*. 2024 Oct 25:e14394. doi: 10.1111/acel.14394. Epub ahead of print. PMID: 39460379.  
Young Finns Study

Schmitz, L. L., Duffie, E., Zhao, W., Ratliff, S. M., Ding, J., Liu, Y., Merkin, S. S., Smith, J. A., & Seeman, T. (2023). Associations of Early-Life Adversity With Later-Life Epigenetic Aging Profiles in the Multi-Ethnic Study of Atherosclerosis. *Am J Epidemiol*, 192(12), 1991-2005. <https://doi.org/10.1093/aje/kwad172>  
MESA

Schmitz, L. L., & Duque, V. (2022). In utero exposure to the Great Depression is reflected in late-life epigenetic aging signatures. *Proc Natl Acad Sci U S A*, 119(46), e2208530119.  
<https://doi.org/10.1073/pnas.2208530119>  
HRS

Schowe, A., Czamara, D., Moenkediek, B., Kuznetsov, D. V., Rohm, T., Instinske, J., Kandler, C., Pahnke, C., Forstner, A., Maj, C., Noethen, M., & Binder, E. B. (2024). Accelerated biological aging during the Covid-19 pandemic – a twinlife epigenetic change satellite study. *Neuroscience Applied*, 3, 104031.  
<https://doi.org/https://doi.org/10.1016/j.nsa.2024.104031>  
TwinLife Epigenetic Change Satellite

Simons, R. L., Ong, M. L., Lei, M. K., Klopack, E., Berg, M., Zhang, Y., Philibert, R., & Beach, S. S. R. (2022). Unstable Childhood, Adult Adversity, and Smoking Accelerate Biological Aging Among Middle-Age African Americans: Similar Findings for GrimAge and PoAm. *J Aging Health*, 34(4-5), 487-498. <https://doi.org/10.1177/08982643211043668>  
FACHS

Uwizeye, G., Rivera, L., Stolrow, H., Christensen, B., Rutherford, J., & Thayer, Z. (2024). Prenatal exposure to genocide accelerates epigenetic aging as measured in second-generation clocks among young adults. *medRxiv*, 2024-10.

## PUBLICATIONS ON MENTAL HEALTH, DRUG & ALCOHOL ABUSE

- Allen, J. P., Danoff, J. S., Costello, M. A., Hunt, G. L., Hellwig, A. F., Krol, K. M., Gregory, S. G., Giamberardino, S. N., Sugden, K., & Connelly, J. J. (2022). Lifetime marijuana use and epigenetic age acceleration: A 17-year prospective examination. *Drug Alcohol Depend*, 233, 109363. <https://doi.org/10.1016/j.drugalcdep.2022.109363>  
South Eastern US cohort
- Apsley, A. T., Ye, Q., Etzel, L., Wolf, S., Hastings, W. J., Mattern, B. C., Siegel, S. R., & Shalev, I. (2023). Biological stability of DNA methylation measurements over varying intervals of time and in the presence of acute stress. *Epigenetics*, 18(1), 2230686.  
<https://doi.org/10.1080/15592294.2023.2230686>  
Texas Twin Study
- Aronoff, J. E., Koning, S. M., Adair, L. S., Lee, N. R., Carba, D. B., Kuzawa, C. W., & McDade, T. W. (2024). Intimate partner violence, depression, and chronic low-grade inflammation among middle-aged women in Cebu, Philippines. *Am J Hum Biol*, e24053.  
<https://doi.org/10.1002/ajhb.24053>  
CEBU
- Beach, S. R. H., Ong, M. L., Gibbons, F. X., Gerrard, M., Lei, M. K., Dawes, K., & Philibert, R. A. (2022). Epigenetic and Proteomic Biomarkers of Elevated Alcohol Use Predict Epigenetic Aging and Cell-Type variation Better Than Self-Report. *Genes (Basel)*, 13(10). <https://doi.org/10.3390/genes13101888>
- Bourassa, K. J., Caspi, A., Brennan, G. M., Hall, K. S., Harrington, H., Houts, R., Kimbrel, N. A., Poulton, R., Ramrakha, S., Taylor, G. A., & Moffitt, T. E. (2023). Which Types of Stress Are Associated With Accelerated Biological Aging? Comparing Perceived Stress, Stressful Life Events, Childhood Adversity, and Posttraumatic Stress Disorder. *Psychosom Med*, 85(5), 389-396. <https://doi.org/10.1097/psy.0000000000001197>  
Dunedin Study
- Bourassa, K. J., Caspi, A., Harrington, H., Houts, R., Poulton, R., Ramrakha, S., & Moffitt, T. E. (2020). Intimate partner violence and lower relationship quality are associated with faster biological aging. *Psychol Aging*, 35(8), 1127-1139. <https://doi.org/10.1037/pag0000581>  
Dunedin Study
- Bourassa, K. J., Garrett, M. E., Caspi, A., Dennis, M., Hall, K. S., Moffitt, T. E., Taylor, G. A., Ashley-Koch, A. E., Beckham, J. C., & Kimbrel, N. A. (2024). Posttraumatic stress disorder, trauma, and accelerated biological aging among post-9/11 veterans. *Transl Psychiatry*, 14(1), 4. <https://doi.org/10.1038/s41398-023-02704-y>  
Veterans Administration Post-Deployment Mental Health Study (PDMH).
- Caro, J. C., Holuka, C., Menta, G., Turner, J. D., Vögele, C., & D'Ambrosio, C. (2023). Children's internalizing behavior development is heterogeneously associated with the pace of epigenetic aging. *Biol Psychol*, 176, 108463.  
<https://doi.org/10.1016/j.biopsych.2022.108463>

- Carreras-Gallo, N., Dwaraka, V. B., Cáceres, A., Smith, R., Mendez, T. L., Went, H., & Gonzalez, J. R. (2023). Impact of tobacco, alcohol, and marijuana on genome-wide DNA methylation and its relationship with hypertension. *Epigenetics*, 18(1), 2214392. <https://doi.org/10.1080/15592294.2023.2214392>  
Trudiagnostic Biobank
- Caspi, A., Shireby, G., Mill, J., Moffitt, T. E., Sugden, K., & Hannon, E. (2023). Accelerated Pace of Aging in Schizophrenia: Five Case-Control Studies. *Biol Psychiatry*. <https://doi.org/10.1016/j.biopsych.2023.10.023>  
Five Case-control Clinical Studies
- Fischer IC, Na PJ, Nagamatsu ST, et al. Posttraumatic Stress Disorder, Obesity, and Accelerated Epigenetic Aging Among US Military Veterans. *JAMA Psychiatry*. 2024;81(12):1276–1277. doi:10.1001/jamapsychiatry.2024.3403
- Goering, M., Moore, A., Barker-Kamps, M., Patki, A., Tiwari, H. K., & Mrug, S. (2024). Adolescent empathy and epigenetic aging in adulthood: Substance use as a mediator. *Developmental Psychology*. Advance online publication. <https://doi.org/10.1037/dev0001893>
- Jung, J., McCartney, D. L., Wagner, J., Yoo, J., Bell, A. S., Mavromatis, L. A., Rosoff, D. B., Hodgkinson, C. A., Sun, H., Schwandt, M., Diazgranados, N., Smith, A. K., Michopoulos, V., Powers, A., Stevens, J., Bradley, B., Fani, N., Walker, R. M., Campbell, A., . . . Lohoff, F. W. (2023). Additive Effects of Stress and Alcohol Exposure on Accelerated Epigenetic Aging in Alcohol Use Disorder. *Biol Psychiatry*, 93(4), 331-341. <https://doi.org/10.1016/j.biopsych.2022.06.036>  
Generation Scotland, Grady Trauma Project
- Langevin, S., Caspi, A., Barnes, J. C., Brennan, G., Poulton, R., Purdy, S. C., Ramrakha, S., Tanksley, P. T., Thorne, P. R., Wilson, G., & Moffitt, T. E. (2022). Life-Course Persistent Antisocial Behavior and Accelerated Biological Aging in a Longitudinal Birth Cohort. *Int J Environ Res Public Health*, 19(21). <https://doi.org/10.3390/ijerph192114402>  
Dunedin Study
- Lima, C. N., Rubinstein, A., Kumar, A., Jha, R., Soyebo, E., Farhan, M., Dwaraka, V., Prestrud, A. A., Smith, R., Walss-Bass, C., Quevedo, J., Soares, J., & Fries, G. (2023). 513. Metabolic and Inflammatory Changes as Biological Mechanisms Underlying the Accelerated Pace of Epigenetic Aging in Bipolar Disorder. *Biological Psychiatry*, 93(9), S301-S302. <https://doi.org/10.1016/j.biopsych.2023.02.753>
- Meier, M. H., Caspi, A., Ambler, A., Hariri, A. R., Harrington, H., Hogan, S., Houts, R., Knott, A. R., Ramrakha, S., Richmond-Rakerd, L. S., Poulton, R., & Moffitt, T. E. (2022). Preparedness for healthy ageing and polysubstance use in long-term cannabis users: a population-representative longitudinal study. *Lancet Healthy Longev*, 3(10), e703-e714. [https://doi.org/10.1016/s2666-7568\(22\)00201-x](https://doi.org/10.1016/s2666-7568(22)00201-x)  
Dunedin Study
- Noguchi, N., Shirai, T., Suda, A., Hattori, S., Miyauchi, M., Okazaki, S., . . . & Hishimoto, A. (2024). Biological aging analysis based on DNA methylation status for social anxiety disorder. *Neuropsychopharmacology Reports*.

- Philibert, R., Lei, M. K., Ong, M. L., & Beach, S. R. (2024). Objective assessments of smoking and drinking outperform clinical phenotypes in predicting variance in epigenetic aging. *Genes*, 15(7), 869.
- PROSAFF, Protecting Strong African American Families
- Roberts, A. L., Ratanatharathorn, A., Chibnik, L., Zhu, Y., Jha, S., Kang, J. H., ... & Koenen, K. C. (2025). No association of posttraumatic stress disorder with epigenetic aging in women at mid-life: A longitudinal cohort study. *Brain, Behavior, and Immunity*, 123, 672-680.
- Nurses Health Study II
- Saarinen, A., Marttila, S., Mishra, P. P., Lyytikäinen, L.-P., Raitoharju, E., Mononen, N., Sormunen, E., Kähönen, M., Raitakari, O., Hietala, J., Keltikangas-Järvinen, L., & Lehtimäki, T. (2024). Polygenic risk for schizophrenia, social dispositions, and pace of epigenetic aging: Results from the Young Finns Study. *Aging Cell*, 23(3), e14052. <https://doi.org/https://doi.org/10.1111/acel.14052>
- Young Finns Study
- Shirai, T., Okazaki, S., Otsuka, I., Miyachi, M., Tanifugi, T., Shindo, R., Okada, S., Minami, H., Horai, T., Mouri, K., & Hishimoto, A. (2024). Accelerated epigenetic aging in alcohol dependence. *Journal of Psychiatric Research*, 173, 175-182. <https://doi.org/https://doi.org/10.1016/j.jpsychires.2024.03.025>
- Suglia Ms, S. F., Clausing, E. S., Shelton, R. C., Conneely, K., Prada-Ortega, D., DeVivo, I., Factor-Litvak, P., Cirillo, P., Baccarelli, A. A., Cohn, B., & Link, B. G. (2024). Cumulative stress across the lifecourse and biological aging in adulthood. *Psychosom Med*. <https://doi.org/10.1097/psy.0000000000001284>
- Child Health and Development Study
- Wang, H., Bakulski, K. M., Blostein, F., Porath, B. R., Dou, J., Tejera, C. H., & Ware, E. B. (2023). Depressive symptoms are associated with DNA methylation age acceleration in a cross-sectional analysis of adults over age 50 in the United States. *medRxiv*. <https://doi.org/10.1101/2023.04.24.23289052>
- HRS
- Webb, E. K., Zannas, A., Linnstaedt, S., McLean, S. A., Ressler, K., & Harnett, N. (2024). 377. Psychological Resilience Modulates the Association Between Neighborhood Disadvantage and Accelerated Epigenetic Aging in Recent Trauma Survivors. *Biological Psychiatry*, 95(10), S254.
- Wertz, J., Caspi, A., Ambler, A., Broadbent, J., Hancox, R. J., Harrington, H., Hogan, S., Houts, R. M., Leung, J. H., Poulton, R., Purdy, S. C., Ramrakha, S., Rasmussen, L. J. H., Richmond-Rakerd, L. S., Thorne, P. R., Wilson, G. A., & Moffitt, T. E. (2021). Association of History of Psychopathology With Accelerated Aging at Midlife. *JAMA Psychiatry*, 78(5), 530-539. <https://doi.org/10.1001/jamapsychiatry.2020.4626>
- Dunedin Study
- Yusupov, N., Dieckmann, L., Erhart, M., Sauer, S., Rex-Haffner, M., Kopf-Beck, J., Brückl, T. M., Czamara, D., & Binder, E. B. (2023). Transdiagnostic evaluation of epigenetic age acceleration and burden of psychiatric disorders. *Neuropsychopharmacology*, 48(9), 1409-1417. <https://doi.org/10.1038/s41386-023-01579-3>

Biological Classification of Mental Disorders study in Germany (BeCOME)

**PUBLICATIONS ON HEALTH BEHAVIORS, LIFESTYLES AND PSYCHOSOCIAL RISKS  
(diet, physical activity, loneliness, sleep, tobacco smoking)**

AUTHOR Remarkable concordance in associations between epigenetic clocks and health behaviors across three countries. Dec 2024.  
HRS, NICOLA, TILDA

Agbor, F., Stoudmire, T., Goodin, B., & Aroke, E. (2024). Stronger Social Support with Friends Correlate with a Slower Pace of Biological Aging. *The Journal of Pain*, 25(4), 43.

Beach, S. R. H., Klopack, E. T., Carter, S. E., Philibert, R. A., Simons, R. L., Gibbons, F. X., Ong, M. L., Gerrard, M., & Lei, M. K. (2022). Do Loneliness and Per Capita Income Combine to Increase the Pace of Biological Aging for Black Adults across Late Middle Age? *Int J Environ Res Public Health*, 19(20). <https://doi.org/10.3390/ijerph192013421>  
The FACHS Study

Beam, C. R., Bakulski, K. M., Zandi, E., Turkheimer, E., Lynch, M., Gold, A. I., ... Davis, D. W. (2024). Epigenome-wide association study of loneliness in a sample of U.S. middle-aged twins. *Epigenetics*, 19(1). <https://doi.org/10.1080/15592294.2024.2427999>  
Louisville Twin Study

Carboneau, M., Li, Y., Prescott, B., Liu, C., Huan, T., Joehanes, R., ... & Ma, J. (2024). Epigenetic Age Mediates the Association of Life's Essential 8 With Cardiovascular Disease and Mortality. *Journal of the American Heart Association*, e032743.  
Framingham Heart Study

Cheng, X., Wei, Y., Wang, R., Jia, C., Zhang, Z., An, J., Li, W., Zhang, J., & He, M. (2023). Associations of essential trace elements with epigenetic aging indicators and the potential mediating role of inflammation. *Redox Biol*, 67, 102910.  
<https://doi.org/10.1016/j.redox.2023.102910>

Cribb, L., Hodge, A.M., Southey, M.C. et al. Dietary factors and DNA methylation-based markers of ageing in 5310 middle-aged and older Australian adults. *GeroScience* (2024).  
<https://doi.org/10.1007/s11357-024-01341-7>  
Melbourne Collaborative Cohort Study

DiMarzio, Karissa , Darlynn M. Rojo-Wissar, Evelyn Hernandez Valencia, Mikayla Ver Pault, Shane Denherder, Adamari Lopez, Jena Lerch, Georgette Metrailler, Sarah M. Merrill, April Highlander, Justin Parent. (2024) Childhood Adversity and Adolescent Epigenetic Age Acceleration: The Role of Adolescent Sleep Health  
medRxiv 2024.09.02.24312939; doi: <https://doi.org/10.1101/2024.09.02.24312939>  
Future of Families and Child Wellbeing Study

Dwaraka, Varun, B., Lucia, A., Natalia, C.-G., Jennifer, L. R., Tayler, H., Aaron, L., Logan, T., Ryan, S., Tavis, L. M., Hannah, W., Emily, R. E., Matthew, M. C., Erica, D. S., Justin, L.

S., & Christopher, D. G. (2023). Unveiling the Epigenetic Impact of Vegan vs. Omnivorous Diets on Aging: Insights from the Twins Nutrition Study (TwiNS). *medRxiv*, 2023.2012.2026.23300543. <https://doi.org/10.1101/2023.12.26.23300543>  
 Twins Nutrition Study (TwiNS)

Deana M Ferreri, Jay T Sutliffe, Nanette V Lopez, Chloe A Sutliffe, Ryan Smith, Natalia Carreras-Gallo, Varun B Dwaraka, Ann Alexis Prestrud, Joel H Fuhrman,(2024). Slower Pace of Epigenetic Aging and Lower Inflammatory Indicators in Females Following a Nutrient-Dense, Plant-Rich Diet Than Those in Females Following the Standard American Diet, Current Developments in Nutrition, Volume 8, Issue 12,2024,104497,ISSN 2475-2991,<https://doi.org/10.1016/j.cdnut.2024.104497>.

Dye, C. K., Wu, H., Jackson, G. L., Kidane, A., Nkambule, R., Lukhele, N. G., Malinga, B. P., Chekenyere, R., El-Sadr, W. M., Baccarelli, A. A., & Harris, T. G. (2024). Epigenetic aging in older people living with HIV in Eswatini: a pilot study of HIV and lifestyle factors and epigenetic aging. *Clinical Epigenetics*, 16(1), 32. <https://doi.org/10.1186/s13148-024-01629-7>

Fancourt, D., Masebo, L., Finn, S., Mak, H. W., & Bu, F. (2024). Does frequency or diversity of leisure activity matter more for epigenetic ageing? Analyses of arts engagement and physical activity in the UK Household Longitudinal Study. *medRxiv*, 2024-11.  
 UK Household Longitudinal Study

Freilich, C. D., Markon, K. E., Cole, S. W., & Krueger, R. F. (2024). Loneliness, epigenetic age acceleration, and chronic health conditions. *Psychol Aging*.  
<https://doi.org/10.1037/pag0000822>  
 MIDUS

Freilich, C. D., Markon, K. E., Mann, F. D., Cole, S. W., & Krueger, R. F. (2024). Associations Between Loneliness, Epigenetic Aging, and Multimorbidity Through Older Adulthood. *The Journals of Gerontology, Series B: Psychological Sciences and Social Sciences*, 79(12), gbae169.

Goering, M., Tiwari, H. K., Patki, A., Espinoza, C. N., Knight, D. C., & Mrug, S. (2024). Examining Health Behaviors as Mechanisms Linking Earlier Pubertal Timing with Accelerated Epigenetic Aging in Late Adolescence. *Journal of Youth and Adolescence*, 1-19.  
 Healthy Passages study in Birmingham (Alabama)

Gonggrijp, M.A.B., van de Weijer, G.A.S., Bijleveld, C.J.H.C. et al. Negative Life Events and Epigenetic Ageing: A Study in the Netherlands Twin Register. *Behav Genet* (2024).  
<https://doi.org/10.1007/s10519-024-10211-z>  
 Netherlands Twin Register

Harris, Kathleen Mullan, Brandt, L., Lauren, G., Chantel, M., Jess, M. M., Aura Ankita, M., Audrey, L. K., & Allison, E. A. (2024). The Sociodemographic and Lifestyle Correlates of Epigenetic Aging in a Nationally Representative U.S. Study of Younger Adults. *bioRxiv*, 2024.2003.2021.585983. <https://doi.org/10.1101/2024.03.21.585983>  
 Add Health

Hillmann, A. R., Dhingra, R., & Reed, R. G. (2023). Positive social factors prospectively predict younger epigenetic age: Findings from the Health and Retirement Study.

*Psychoneuroendocrinology*, 148, 105988.

<https://doi.org/10.1016/j.psyneuen.2022.105988>

HRS

Kankaanpää, A., Tolvanen, A., Heikkinen, A., Kaprio, J., Ollikainen, M., & Sillanpää, E. (2022). The role of adolescent lifestyle habits in biological aging: A prospective twin study. *eLife*, 11. <https://doi.org/10.7554/eLife.80729>

FinnTwin12

Kankaanpää, A., Tolvanen, A., Joensuu, L., Waller, K., Heikkinen, A., Kaprio, J., Ollikainen, M., & Sillanpää, E. (2023). The associations of long-term physical activity in adulthood with later biological ageing and all-cause mortality - a prospective twin study. *medRxiv*.

<https://doi.org/10.1101/2023.06.02.23290916>

Older Finnish Twin Cohort

Ke, T. M., Lophatananon, A., & Muir, K. R. (2024). Exploring the Relationships between Lifestyle Patterns and Epigenetic Biological Age Measures in Men. *Biomedicines*, 12(9), 1985.

Kim DJ, Kang JH, Kim J-W, Kim Sb, Lee YK, Cheon MJ and Lee B-C (2024) Assessing the utility of epigenetic clocks for health prediction in South Korean. *Front. Aging* 5:1493406.

doi: 10.3389/fragi.2024.1493406

Korea Association Resource (KARE) of the Korean Genome and Epidemiology Study (KoGES) Health Examinees (HEXA) of the KoGES

Kim, E. S., Nakamura, J. S., Strecher, V. J., & Cole, S. W. (2023). Reduced Epigenetic Age in Older Adults With High Sense of Purpose in Life. *J Gerontol A Biol Sci Med Sci*, 78(7), 1092-1099. <https://doi.org/10.1093/gerona/glad092>

HRS

Kim, Y., Huan, T., Joehanes, R., McKeown, N. M., Horvath, S., Levy, D., & Ma, J. (2022).

Higher diet quality relates to decelerated epigenetic aging. *Am J Clin Nutr*, 115(1), 163-170. <https://doi.org/10.1093/ajcn/nqab201>

The Framingham Offspring Study.

Kim, S., Halvorsen, C., Potter, C., & Faul, J. (2025). Does volunteering reduce epigenetic age acceleration among retired and working older adults? Results from the Health and Retirement Study. *Social Science & Medicine*, 364, 117501.

HRS

Klopack, ET, JE Carroll, SW Cole, TE Seeman, EM Crimmins (2022). Lifetime exposure to smoking, epigenetic aging, and morbidity and mortality in older adults. *Clinical Epigenetics* 14 (1), 72

HRS

- Li, D. L., Hodge, A. M., Cribb, L., Southey, M. C., Giles, G. G., Milne, R. L., & Dugué, P. A. (2024). Body size, diet quality, and epigenetic ageing: cross-sectional and longitudinal analyses. *J Gerontol A Biol Sci Med Sci*. <https://doi.org/10.1093/gerona/glae026>
- Loh, K. P., Sanapala, C., Jensen-Battaglia, M., Rana, A., Sohn, M. B., Watson, E., Gilmore, N., Klepin, H. D., Mendler, J. H., Liesveld, J., Huselton, E., LoCastro, M., Susiarjo, M., Netherby-Winslow, C., Williams, A. M., Mustian, K., Vertino, P., & Janelsins, M. C. (2023). Exercise and epigenetic ages in older adults with myeloid malignancies. *Eur J Med Res*, 28(1), 180. <https://doi.org/10.1186/s40001-023-01145-z>
- Michels, K. B., & Binder, A. M. (2024). Impact of folic acid supplementation on the epigenetic profile in healthy unfortified individuals - a randomized intervention trial. *Epigenetics*, 19(1), 2293410. <https://doi.org/10.1080/15592294.2023.2293410>
- Nakamura, J. S., Kwok, C., Huang, A., Strecher, V. J., Kim, E. S., & Cole, S. W. (2023). Reduced epigenetic age in older adults who volunteer. *Psychoneuroendocrinology*, 148, 106000. <https://doi.org/10.1016/j.psyneuen.2022.106000>  
HRS
- Norooz, R., Rudnicka, J., Pisarek, A., Wysocka, B., Masny, A., Boroń, M., Migacz-Gruszka, K., Pruszkowska-Przybylska, P., Kobus, M., Lisman, D., Zielińska, G., Iljin, A., Wiktorska, J., Michalczyk, M., Kaczka, P., Krzysztofik, M., Sitek, A., Ossowski, A., Spólnicka, M., . . . Pośpiech, E. (2024). Analysis of epigenetic clocks links yoga, sleep, education, reduced meat intake, coffee, and a SOCS2 gene variant to slower epigenetic aging. *Geroscience*, 46(2), 2583-2604. <https://doi.org/10.1007/s11357-023-01029-4>
- Ravi, Suvi et al. (2024). Suboptimal dietary patterns are associated with accelerated biological aging in young adulthood: a study with twins, 2024 , Clinical Nutrition, Finntwin12
- Rentscher, K. E., Klopack, E. T., Crimmins, E. M., Seeman, T. E., Cole, S. W., & Carroll, J. E. (2023). Social relationships and epigenetic aging in older adulthood: Results from the Health and Retirement Study. *Brain Behav Immun*, 114, 349-359. <https://doi.org/10.1016/j.bbi.2023.09.001>  
HRS
- Reynolds, L. M., Houston, D. K., Skiba, M. B., Whitsel, E. A., Stewart, J. D., Li, Y., Zannas, A. S., Assimes, T. L., Horvath, S., Bhatti, P., Baccarelli, A. A., Tooze, J. A., & Vitolins, M. Z. (2024). Diet Quality and Epigenetic Aging in the Women's Health Initiative. *J Acad Nutr Diet*. <https://doi.org/10.1016/j.jand.2024.01.002>  
Women's Health Initiative
- Simons, R. L., Ong, M. L., Lei, M. K., Beach, S. R. H., Zhang, Y., Philibert, R., & Mielke, M. M. (2023). Changes in Loneliness, BDNF, and Biological Aging Predict Trajectories in a Blood-Based Epigenetic Measure of Cortical Aging: A Study of Older Black Americans. *Genes (Basel)*, 14(4). <https://doi.org/10.3390/genes14040842>  
FACHS
- Simons, R. L., Ong, M. L., Lei, M. K., Klopach, E., Berg, M., Zhang, Y., Philibert, R., Gibbons, F. X., & Beach, S. R. H. (2022). Shifts in lifestyle and socioeconomic circumstances predict

change-for better or worse-in speed of epigenetic aging: A study of middle-aged black women. *Soc Sci Med*, 307, 115175. <https://doi.org/10.1016/j.socscimed.2022.115175>  
FACHS

Simons, R. L., Ong, M. L., Lei, M. K., Klopack, E., Berg, M., Zhang, Y., Philibert, R., & Beach, S. S. R. (2022). Unstable Childhood, Adult Adversity, and Smoking Accelerate Biological Aging Among Middle-Age African Americans: Similar Findings for GrimAge and PoAm. *J Aging Health*, 34(4-5), 487-498. <https://doi.org/10.1177/08982643211043668>  
FACHS

Streck, B., Hyun, G., Armstrong, G. T., Belsky, D. W., Ehrhardt, M. J., Hudson, M., Green, P., Robison, L. L., Tonorezos, E. S., Yasui, Y., Wang, Z., Wilson, C. L., Brinkman, T. M., Ness, K. K., & Guida, J. L. (2023). Associations between psychosocial and lifestyle factors and biological age acceleration among long-term survivors of childhood cancer: A prospective study from the St. Jude Lifetime Cohort. *Journal of Clinical Oncology*, 41(16\_suppl), 10049-10049. [https://doi.org/10.1200/JCO.2023.41.16\\_suppl.10049](https://doi.org/10.1200/JCO.2023.41.16_suppl.10049)  
St. Jude Lifetime Cohort

Thomas, A., Ryan, C. P., Caspi, A., Moffitt, T. E., Sugden, K., Zhou, J., Belsky, D. W., & Gu, Y. (2023). Diet, pace of biological aging, and risk of dementia in the Framingham Heart Study. *medRxiv*. <https://doi.org/10.1101/2023.05.24.23290474>  
Framingham Offspring Study

Wang, W., Dearman, A., Bao, Y., & Kumari, M. (2023). Partnership status and positive DNA methylation age acceleration across the adult lifespan in the UK. *SSM Popul Health*, 24, 101551. <https://doi.org/10.1016/j.ssmph.2023.101551>  
Understanding Society

Waziry, R., Ryan, C. P., Corcoran, D. L., Huffman, K. M., Kobor, M. S., Kothari, M., Graf, G. H., Kraus, V. B., Kraus, W. E., Lin, D. T. S., Pieper, C. F., Ramaker, M. E., Bhapkar, M., Das, S. K., Ferrucci, L., Hastings, W. J., Kebbe, M., Parker, D. C., Racette, S. B., . . . Belsky, D. W. (2023). Effect of long-term caloric restriction on DNA methylation measures of biological aging in healthy adults from the CALERIE trial. *Nat Aging*, 3(3), 248-257. <https://doi.org/10.1038/s43587-022-00357-y>  
CALERIE TRIAL

Willem, Y. E., deSteiguer, A., Tanksley, P. T., Vinnik, L., Främke, D., Okbay, A., Richter, D., Wagner, G. G., Hertwig, R., Koellinger, P., Tucker-Drob, E. M., Harden, K. P., & Raffington, L. (2023). Self-control is associated with health-relevant disparities in buccal DNA-methylation measures of biological aging in older adults. *medRxiv*.  
<https://doi.org/10.1101/2023.08.30.23294816>  
German Socioeconomic Panel Study (SOEP-Gene)

Yu, Y. L. (2023). Current Marital Status and Epigenetic Clocks Among Older Adults in the United States: Evidence From the Health and Retirement Study. *J Aging Health*, 35(1-2), 71-82. <https://doi.org/10.1177/08982643221104928>  
HRS

Zhang, A., Zhang, Y., Meng, Y., Ji, Q., Ye, M., Zhou, L., Liu, M., Yi, C., Karlsson, I. K., Fang, F., Hägg, S., & Zhan, Y. (2024). Associations between psychological resilience and

epigenetic clocks in the health and retirement study. *Geroscience*, 46(1), 961-968.  
<https://doi.org/10.1007/s11357-023-00940-0>  
 HRS

Zheng, H. T., Li, D. L., Lou, M. W., Hodge, A. M., Southey, M. C., Giles, G. G., ... & Dugué, P. A. (2024). Physical activity and DNA methylation-based markers of ageing in 6208 middle-aged and older Australians: cross-sectional and longitudinal analyses. *GeroScience*, 1-12.

Melbourne Collaborative Cohort Study

### PUBLICATIONS ON TOXIC EXPOSURES

Boyer, K., Domingo-Relloso, A., Jiang, E., Haack, K., Goessler, W., Zhang, Y., Umans, J. G., Belsky, D. W., Cole, S. A., Navas-Acien, A., & Kupsco, A. (2023). Metal mixtures and DNA methylation measures of biological aging in American Indian populations. *Environ Int*, 178, 108064. <https://doi.org/10.1016/j.envint.2023.108064>  
 The Strong Heart Study of Native Americans

Chang, C. J., O'Brien, K. M., Kresovich, J. K., Nwanaji-Enwerem, J. C., Xu, Z., Gaston, S. A., ... & White, A. J. (2024). Associations between use of chemical hair products and epigenetic age: Findings from the Sister Study. *Environmental Epidemiology*, 8(3), e311. Sister Study

Gao, X., Huang, J., Cardenas, A., Zhao, Y., Sun, Y., Wang, J., Xue, L., Baccarelli, A. A., Guo, X., Zhang, L., & Wu, S. (2022). Short-Term Exposure of PM(2.5) and Epigenetic Aging: A Quasi-Experimental Study. *Environ Sci Technol*, 56(20), 14690-14700. <https://doi.org/10.1021/acs.est.2c05534>

Jiang, E. X., Domingo-Relloso, A., Abuawad, A., Haack, K., Tellez-Plaza, M., Fallin, M. D., Umans, J. G., Best, L. G., Zhang, Y., Kupsco, A., Belsky, D. W., Cole, S. A., & Navas-Acien, A. (2023). Arsenic Exposure and Epigenetic Aging: The Association with Cardiovascular Disease and All-Cause Mortality in the Strong Heart Study. *Environ Health Perspect*, 131(12), 127016. <https://doi.org/10.1289/ehp11981>  
 The Strong Heart Study of Native Americans

Koenigsberg, S. H., Chang, C. J., Ish, J., Xu, Z., Kresovich, J. K., Lawrence, K. G., Kaufman, J. D., Sandler, D. P., Taylor, J. A., & White, A. J. (2023). Air pollution and epigenetic aging among Black and White women in the US. *Environ Int*, 181, 108270. <https://doi.org/10.1016/j.envint.2023.108270>  
 KORA (Cooperative Health Research in the Region Augsburg study), NAS (Normative Aging Study)

Lucia, R. M., Huang, W. L., Pathak, K. V., McGilvrey, M., David-Dirgo, V., Alvarez, A., Goodman, D., Masunaka, I., Odegaard, A. O., Ziogas, A., Pirrotte, P., Norden-Krichmar, T. M., & Park, H. L. (2022). Association of Glyphosate Exposure with Blood DNA Methylation in a Cross-Sectional Study of Postmenopausal Women. *Environ Health Perspect*, 130(4), 47001. <https://doi.org/10.1289/ehp10174>

Schmidt, S. (2024). Marking Time: Epigenetic Aging May Partially Explain the Arsenic-Cardiovascular Disease Link. *Environmental Health Perspectives*, 132(2), 024001.

Shi, W., Tang, S., Fang, J., Cao, Y., Chen, C., Li, T., Gao, X., & Shi, X. (2022). Epigenetic age stratifies the risk of blood pressure elevation related to short-term PM(2.5) exposure in older adults. *Environ Res*, 212(Pt D), 113507. <https://doi.org/10.1016/j.envres.2022.113507>

Wu, Y., Xu, R., Li, S., Wen, B., Southey, M. C., Dugue, P. A., ... & Guo, Y. (2025). Association between wildfire-related PM<sub>2.5</sub> and epigenetic aging: A twin and family study in Australia. *Journal of Hazardous Materials*, 481, 136486.

Yannatos, I., Stites, S., Brown, R. T., & McMillan, C. T. (2023). Contributions of neighborhood social environment and air pollution exposure to Black-White disparities in epigenetic aging. *PLoS One*, 18(7), e0287112. <https://doi.org/10.1371/journal.pone.0287112>  
HRS

### **EXPERIMENTAL INTERVENTION STUDIES and Natural Experiments**

Borrus, D. S., Sehgal, R., Armstrong, J. F., Kasamoto, J., Gonzalez, J., & Higgins-Chen, A. T. (2024). When to Trust Epigenetic Clocks: Avoiding False Positives in Aging Interventions. *bioRxiv*, 2024-10.

Carroll, Jude (GSA symposium submission 2023): **Intervention** delivered to older adults with insomnia, is associated with slowed DunedinPACE from baseline to 18-24 month follow-up.

Corley, Michael J., Alina P. S. Pang, Cecilia M. Shikuma, Lishomwa C. Ndhlovu (2023). Cell-type specific impact of **metformin** on monocyte epigenetic age reversal in virally suppressed older people living with HIV. *Aging Cell*. <https://doi.org/10.1111/acel.13926>

Dawson, Kristin, Athena May Jean M. Carangan, Jessica Klunder, Natalia Carreras-Gallo, Raghav Sehgal, Samantha Megilligan, Benjamin C. Askins, Nicole Perkins, Tavis L. Mendel, Ryan Smith, Matthew Dawson, Michael Mallin, Albert T. Higgins-Chen, Varun B. Dwaraka. (2024) Ketamine treatment effects on DNA methylation and Epigenetic Biomarkers of aging, *medArXiv*

Drouard, G., Suhonen, S., Heikkinen, A., Wang, Z., Kaprio, J., & Ollikainen, M. (2024). Multi-omic associations of epigenetic age acceleration are heterogeneously shaped by genetic and environmental influences. *medRxiv*, 2024-06.

Dwaraka, Varun B., Lucia Aronica, Natalia Carreras-Gallo, Jennifer L Robinson, Tayler Hennings, Aaron Lin, Logan Turner, Ryan Smith, Tavis L. Mendez, Hannah Went, Emily R. Ebel, Matthew M. (in review). Unveiling the Epigenetic Impact of **Vegan vs. Omnivorous Diets** on Aging: Insights from the Twins Nutrition Study (TwINS). <https://www.medrxiv.org/content/medrxiv/early/2023/12/29/2023.12.26.23300543.full.pdf>

Lee, E., Carreras-Gallo, N., Lopez, L., Turner, L., Lin, A., Mendez, T. L., Went, H., Tomusiak, A., Verdin, E., Corley, M., Ndhlovu, L., Smith, R., & Dwaraka, V. B. (2024). Exploring the effects of Dasatinib, Quercetin, and Fisetin on DNA methylation clocks: a longitudinal

study on senolytic interventions. *Aging (Albany NY)*, 16(4), 3088-3106.  
<https://doi.org/10.18632/aging.205581>

Holland, P., Istre, M., Ali, M. M., Gedde-Dahl, T., Buechner, J., Wildhagen, M., Brunvoll, S. H., Horvath, S., Matsuyama, S., Dahl, J. A., Stölzel, F., & Søraas, A. (2024). Epigenetic aging of human blood cells is influenced by the age of the host body. *Aging Cell*, e14112. <https://doi.org/10.1111/ace.14112>

Merrill SM, Hogan C, Bozack AK, et al. Telehealth Parenting Program and Salivary Epigenetic Biomarkers in Preschool Children With Developmental Delay: NIMHD Social Epigenomics Program. *JAMA Netw Open*.2024;7(7):e2424815. doi:10.1001/jamanetworkopen.2024.24815

Michels, Karin B. & Alexandra M. Binder (2024) Impact of **folic acid supplementation** on the epigenetic profile in healthy unfortified individuals – a randomized intervention trial, *Epigenetics*, 19:1, DOI: [10.1080/15592294.2023.2293410](https://doi.org/10.1080/15592294.2023.2293410)

Nannini, D. R., Cortese, R., VonTungeln, C., & Hildebrandt, G. C. (2024). Chemotherapy-induced acceleration of DNA methylation-based biological age in breast cancer. *Epigenetics*, 19(1), 2360160.

Poganik, Jesse R., Bohan Zhang, Gurpreet S. Baht, Csaba Kerepesi, Sun Hee Yim, Ake T. Lu, Amin Haghani, Tong Gong, Anna M. Hedman, Ellika Andolf, Göran Pershagen, Catarina Almqvist, James P. White, Steve Horvath, Vadim N. Gladyshev, Biological age is increased by **stress**, and restored upon **recovery from stress**, *Cell Metabolism* 2023.  
 Three Clinical Datasets

Sehgal, Raghav, Daniel Borrus, Jessica Kasamato, Jenel F. Armstrong, John Gonzalez, Yaroslav Markov, Ahana Priyanka, Ryan Smith, Natàlia Carreras, Varun B. Dwaraka, Albert Higgins-Chen, DNAm aging biomarkers are responsive: Insights from 51 longevity interventional studies in humans, BioRxiv, doi: <https://doi.org/10.1101/2024.10.22.619522>

Sullivan, J., Nicholson, T., Hazeldine, J. et al. Accelerated epigenetic ageing after burn injury. *GeroScience* (2025). <https://doi.org/10.1007/s11357-024-01433-4>

Waziry, R., Ryan, C. P., Corcoran, D. L., Huffman, K. M., Kobor, M. S., Kothari, M., Graf, G. H., Kraus, V. B., Kraus, W. E., Lin, D. T. S., Pieper, C. F., Ramaker, M. E., Bhapkar, M., Das, S. K., Ferrucci, L., Hastings, W. J., Kebbe, M., Parker, D. C., Racette, S. B., . . . Belsky, D. W. (2023). Effect of long-term caloric restriction on DNA methylation measures of biological aging in healthy adults from the CALERIE trial. *Nat Aging*, 3(3), 248-257. <https://doi.org/10.1038/s43587-022-00357-y>  
 CALERIE

Yaskolka Meir, A., Keller, M., Hoffmann, A., Rinott, E., Tsaban, G., Kaplan, A., Zelicha, H., Hagemann, T., Ceglarek, U., Isermann, B., Shelef, I., Blüher, M., Stumvoll, M., Li, J., Haange, S. B., Engelmann, B., Rolle-Kampczyk, U., von Bergen, M., Hu, F. B., . . . Shai, I. (2023). The effect of polyphenols on DNA methylation-assessed biological age

attenuation: the DIRECT PLUS randomized controlled trial. *BMC Med*, 21(1), 364. <https://doi.org/10.1186/s12916-023-03067-3>

### **PUBLICATIONS ON MECHANISMS AFFECTING CLOCKS**

Apsley, A. T., Ye, Q., Etzel, L., Wolf, S., Hastings, W. J., Mattern, B. C., ... Shalev, I. (2023). Biological stability of DNA methylation measurements over varying intervals of time and in the presence of acute stress. *Epigenetics*, 18(1). <https://doi.org/10.1080/15592294.2023.2230686> <https://www.tandfonline.com/doi/full/10.1080/15592294.2023.2230686>

Asefa, N., Li, Z., Guðmundsdóttir, V., Gudnason, V., & Launer, L. (2024). Five-Year Change in Biological Aging: Insights From the Age Gene-Environment Susceptibility-Reykjavik Study. *Circulation*, 149(Suppl\_1), A47-A47.

Age Gene/Environment Susceptibility-Reykjavik Study (AGES-RS)

deSteiguer, A. J., Raffington, L., Sabhlok, A., Tanksley, P., Tucker-Drob, E. M., & Harden, K. P. (2023). Stability of DNA-Methylation Profiles of Biological Aging in Children and Adolescents. *bioRxiv*. <https://doi.org/10.1101/2023.10.30.564766>  
Texas Twin St

Dowrey, T. W., Cranston, S. F., Skvir, N., Lok, Y., Gould, B., Petrowitz, B., Villar, D., Shan, J., James, M., Dodge, M., Belkina, A. C., Giadone, R. M., Sebastiani, P., Perls, T. T., Andersen, S. L., & Murphy, G. J. (2024). A longevity-specific bank of induced pluripotent stem cells from centenarians and their offspring. *Aging Cell*, e14351

Drewelies, J., Homann, J., Vetter, V. M., Duezel, S., Kühn, S., Deecke, L., ... & Gerstorf, D. (2024). There are multiple clocks that time us: Cross-sectional and longitudinal associations among 14 alternative indicators of age and aging. *The Journals of Gerontology, Series A: Biological Sciences and Medical Sciences*, glae244.

Hong, X., Cao, H., Cao, W., Lv, J., Yu, C., Huang, T., ... & Li, L. (2024). Trends of genetic contributions on epigenetic clocks and related methylation sites with aging: A population-based adult twin study. *Aging Cell*, e14403.

Kuznetsov, D. V., Liu, Y., Schowe, A. M., Czamara, D., Instinske, J., Pahnke, C. K., ... & Moenckediek, B. (2024). Age-Associated Genetic and Environmental Contributions to Epigenetic Aging Across Adolescence and Emerging Adulthood. *bioRxiv*, 2024-06.

Marttila, S., Rajić, S., Ciantar, J., Mak, J. K., Junntila, I. S., Kummola, L., ... & Kananen, L. (2024). Biological aging of different blood cell types. *GeroScience*, 1-18.

Miao, K., Shunkai, L., Tao Huang, D., Sun, C. L., Yuanjie Pang, R. H., Zengchang Pang, M. Y., Hua, W., Xianping Wu, Y. L., & Wenjing Gao, L. L. Five years of change in adult twins: longitudinal changes of genetic and environmental influence on epigenetic clocks. *BMC Med* 22, 289 (2024). <https://doi.org/10.1186/s12916-024-03511-y>  
Chinese National Twin Registry

Mulder, Rosa H. et al. (2024) Characterising developmental dynamics of adult epigenetic clock sites. *eBioMedicine*, Volume 109, 105425

Ryan, C. P., Lee, N. R., Carba, D. B., MacIsaac, J. L., Lin, D. S., Atashzay, P., Belsky, D. W., Kober, M. S., McDade, T. W., & Kuzawa, C. W. (In Press). Pregnancy is linked to faster epigenetic aging in young women. *Proceedings of the National Academy of Sciences of the USA*.

Schmid, H., Vetter, V. M., Homann, J., Bahr, V., Lill, C. M., Regitz-Zagrosek, V., ... & Demuth, I. (2024). Cross-sectional and Longitudinal Relationship between Sex Hormones and Six Epigenetic Clocks in Older Adults: Results of the Berlin Aging Study II (BASE-II). *medRxiv*, 2024-11.

Ying, K., Liu, H., Tarkhov, A. E., Sadler, M. C., Lu, A. T., Moqri, M., Horvath, S., Katalik, Z., Shen, X., & Gladyshev, V. N. (2024). Causality-enriched epigenetic age uncouples damage and adaptation. *Nat Aging*, 4(2), 231-246. <https://doi.org/10.1038/s43587-023-00557-0>

Zhang, Z., Reynolds, S. R., Stolrow, H. G., Chen, J. Q., Christensen, B. C., & Salas, L. A. (2024). Deciphering the role of immune cell composition in epigenetic age acceleration: Insights from cell-type deconvolution applied to human blood epigenetic clocks. *Aging Cell*, 23(3), e14071. <https://doi.org/10.1111/acel.14071>

## LITERATURE REVIEWS

Belsky, D. W., & Baccarelli, A. A. (2023). To promote healthy aging, focus on the environment. *Nat Aging*, 3(11), 1334-1344. <https://doi.org/10.1038/s43587-023-00518-7>

Bourassa, K.J., Sbarra, D.A. Trauma, adversity, and biological aging: behavioral mechanisms relevant to treatment and theory. *Transl Psychiatry* 14, 285 (2024).  
<https://doi.org/10.1038/s41398-024-03004-9>

Espinoza, S. E., Khosla, S., Baur, J. A., de Cabo, R., & Musi, N. (2023). Drugs Targeting Mechanisms of Aging to Delay Age-Related Disease and Promote Healthspan: Proceedings of a National Institute on Aging Workshop. *The Journals of Gerontology: Series A*, 78(Supplement\_1), 53-60. <https://doi.org/10.1093/gerona/glad034>

Gaylord, A., Cohen, A., & Kupsco, A. (2023). Biomarkers of aging through the life course: A Recent Literature Update. *Curr Opin Epidemiol Public Health*, 2(2), 7-17.  
<https://doi.org/10.1097/pxh.0000000000000018>

Herzog, Chiara, Ludger J. E. Goeminne<sup>2,36</sup>, Jesse R. Poganik<sup>2,36</sup>, Nir Barzilai<sup>3</sup>, Daniel W. Belsky<sup>4</sup>, Joe Betts-LaCroix<sup>5</sup>, Brian H. Chen<sup>6</sup>, Michelle Chen<sup>7</sup>, Alan A. Cohen<sup>8</sup>, Steven Cummings<sup>9,10</sup>, Peter O. Fedichev<sup>11</sup>, Luigi Ferrucci<sup>12</sup>, Alexander Fleming<sup>13</sup>, Kristen Fortney<sup>14</sup>, David Furman<sup>15,16,17</sup>, Vera Gorbunova<sup>18</sup>, Albert Higgins-Chen<sup>19</sup>, Lee Hood<sup>15,20</sup>, Steve Horvath<sup>21</sup>, Jamie N. Justice<sup>22,23</sup>, Douglas Kiel<sup>24,25</sup>, George Kuchel<sup>26</sup>, Jessica Lasky-Su<sup>27</sup>, Nathan LeBrasseur<sup>28</sup>, Andrea B. Maier<sup>29,30</sup>, Birgit

Schilling15, Vittorio Sebastian031, P. Eline Slagboom32, Michael P. Snyder33, Eric Verdin15, Martin Widschwendter1,34,35, Alex Zhavoronkov7, Mahdi Moqri2,33  & Vadim N. Gladyshev2  (2024). Challenges and recommendations for the translation of biomarkers of aging. *Nature Ageing*

Justice, J. N., & Kritchevsky, S. B. (2020). Putting epigenetic biomarkers to the test for clinical trials. *eLife*, 9. <https://doi.org/10.7554/eLife.58592>

Kejun, Y., Alexander, T., Alexandre, T., Hanna, L., Mahdi, M., Csaba, K., & Vadim, N. G. (2023). <em>ClockBase</em>; a comprehensive platform for biological age profiling in human and mouse. *bioRxiv*, 2023.2002.2028.530532. <https://doi.org/10.1101/2023.02.28.530532>

Cynthia D.J. Kusters and Steve Horvath (2025) Quantification of Epigenetic Aging in Public Health, Annual Review of Public Health, Vol. 46 <https://doi.org/10.1146/annurev-publhealth-060222-015657>

Leviton, Alan, Olaf Dammann, Anup D. Patel, and Tobias Loddenkemper (2024) Biologic Correlates and Consequences of the Social Determinants of Health and Disease. Perspectives in Biology and Medicine, Johns Hopkins University Press, Volume 67, Number 3, Summer 2024, pp. 305-324, 10.1353/pbm.2024.a936212

Martínez-Magaña, J. J., Hurtado-Soriano, J., Rivero-Segura, N. A., Montalvo-Ortiz, J. L., García-delaTorre, P., Becerril-Rojas, K., & Gomez-Verjan, J. C. (2024). Towards a novel frontier in the use of epigenetic clocks in epidemiology. *Archives of Medical Research*, 55(5), 103033.

Min, M., Egli, C., Dulai, A. S., & Sivamani, R. K. (2024). Critical review of aging clocks and factors that may influence the pace of aging. *Frontiers in Aging*, 5, 1487260.

Moqri, M., Herzog, C., Poganik, J. R., Justice, J., Belsky, D. W., Higgins-Chen, A., Moskalev, A., Fuellen, G., Cohen, A. A., Bautmans, I., Widschwendter, M., Ding, J., Fleming, A., Mannick, J., Han, J. J., Zhavoronkov, A., Barzilai, N., Kaeberlein, M., Cummings, S., . . . Gladyshev, V. N. (2023). Biomarkers of aging for the identification and evaluation of longevity interventions. *Cell*, 186(18), 3758-3775. <https://doi.org/10.1016/j.cell.2023.08.003>

Moqri, M., Herzog, C., Poganik, J. R., Ying, K., Justice, J. N., Belsky, D. W., Higgins-Chen, A. T., Chen, B. H., Cohen, A. A., Fuellen, G., Hägg, S., Marioni, R. E., Widschwendter, M., Fortney, K., Fedichev, P. O., Zhavoronkov, A., Barzilai, N., Lasky-Su, J., Kiel, D. P., . . . Ferrucci, L. (2024). Validation of biomarkers of aging. *Nature Medicine*, 30(2), 360-372. <https://doi.org/10.1038/s41591-023-02784-9>

Noterman, D. A. (2021). Growing Old Before Their Time: Measuring Aging. *Pediatrics*, 147(6). <https://doi.org/10.1542/peds.2021-049939>

Rutledge, J., Oh, H., & Wyss-Coray, T. (2022). Measuring biological age using omics data. *Nat Rev Genet*, 23(12), 715-727. <https://doi.org/10.1038/s41576-022-00511-7>

Ubaida-Mohien, C., Tanaka, T., Tian, Q., Moore, Z., Moaddel, R., Basisty, N., Simonsick, E. M., & Ferrucci, L. (2023). Blood Biomarkers for Healthy Aging. *Gerontology*, 69(10), 1167-1174. <https://doi.org/10.1159/000530795>

Ying, Kejun, Seth Paulson, Alec Eames, Alexander Tyshkovskiy, Siyuan Li, Martin Perez-Guevara, Mehrnoosh Emamifar, Maximiliano Casas Martínez, Dayoon Kwon, Anna Kosheleva, Michael P. Snyder, Dane Gobel, Chiara Herzog, Jesse R. Paganik, Biomarker of Aging Consortium, Mahdi Moqri, Vadim N. Gladyshev. (2024 preprint) A Unified Framework for Systematic Curation and Evaluation of Aging Biomarkers.

<https://www.researchsquare.com/article/rs-4481437/v1.pdf>

Normative Aging Study, MassGeneral Brigham dataset

Zhou, A., Wu, Z., Zaw Phyo, A. Z., Torres, D., Vishwanath, S., & Ryan, J. (2022). Epigenetic aging as a biomarker of dementia and related outcomes: a systematic review. *Epigenomics*, 14(18), 1125-1138. <https://doi.org/10.2217/epi-2022-0209>

## SUPPLEMENTAL MATERIALS:

Idan Shalev reports that most epigenetic sites are unstable, except those in Dunedin PACE (and in PC clocks). <https://www.tandfonline.com/doi/full/10.1080/15592294.2023.2230686>

**Clockbase:** Many studies that have derived DunedinPACE have joined ClockBASE, an open access computational biology data repository: <http://gladyshevlab.org:3838/ClockBase/>.

Ying, K., Tyshkovskiy, A., Trapp, A., Liu, H., Moqri, M., Kerepesi, C., & Gladyshev, V. N. (2023). ClockBase: a comprehensive platform for biological age profiling in human and mouse. *bioRxiv*, 2023-02.

**Biolearn** <https://bio-learn.github.io/>, is an open source library to help with calculating any composite biomarker of aging on any molecular data. This library will standardize all the public biomarker formulations, including the collection of epigenetic clocks standardized by Morgan Levine/Albert Higgins-Chen's ([methylCIPHER](#)) and Vadim Gladyshev's ([ClockBase](#)) groups as well as the blood biomarkers standardized by Daniel Belsky's group ([BioAge](#)). At the same time, Biolearn will harmonize public and private molecular datasets commonly used for biomarker validation. Biolearn is scientifically and financially supported by [Biomarkers of Aging Consortium](#) and [Methuselah Foundation](#).

Ying, Kejun, Seth Paulson, Martin Perez-Guevara, Mehrnoosh Emamifar, Maximiliano Casas Martínez, Dayoon Kwon, Jesse R. Paganik, Mahdi Moqri, Vadim N. Gladyshev, (2023) Biolearn, an open-source library for biomarkers of aging. *BioRxiv*, 4 Dec 2023 <https://www.biorxiv.org/content/biorxiv/early/2023/12/04/2023.12.02.569722.full.pdf>

Ying, Kejun, Seth Paulson, Alec Eames, Alexander Tyshkovskiy, Siyuan Li, Martin Perez-Guevara, Mehrnoosh Emamifar, Maximiliano Casas Martínez, Dayoon Kwon, Anna Kosheleva, Michael P. Snyder, Dane Gobel, Chiara Herzog, Jesse R. Paganik, Biomarker of Aging Consortium, Mahdi Moqri, Vadim N. Gladyshev. (2024 preprint) A Unified Framework for

Systematic Curation and Evaluation of Aging Biomarkers.  
<https://www.researchsquare.com/article/rs-4481437/v1.pdf>  
 Normative Aging Study, MassGeneral Brigham dataset

Rayevskiy, S., Le, Q., Nguyen, J., Chen, S. Q., & Castellani, C. A. (2024).  
 EpigeneticAgePipeline: an R package for comprehensive assessment of epigenetic age metrics from methylation microarrays. *bioRxiv*, 10.

Python aging biomarker access: GitHub is here <https://github.com/rsinghlab/pyaging>.

The preprint is here <https://www.biorxiv.org/content/10.1101/2023.11.28.569069v1>.

The designer: <https://www.linkedin.com/in/lucas-paulo-de-lima-camillo-80706b144/?originalSubdomain=uk>, works for Shift Biosciences --  
<https://www.shiftbioscience.com/>

**TruDiagnostic report** (Licensed Feb 2021).

<https://drive.google.com/file/d/1ZXdQhc8io9InMypibr6C-MI1BC-hPugy/view>

**Everything Epigenetics podcast by Hannah Went:**

<https://everythingepigenetics.com/episode/dr-terrie-moffitt-how-fast-are-you-aging-really-dunedinpace/>

**Bryan Johnson's Blueprint** for slowing aging and **Rejuvenation Olympics** lederboard features DunedinPACE.

<https://www.lifespan.io/news/bryan-johnsons-race-against-time/>  
<https://blueprint.bryanjohnson.co/>  
<https://rejuvenationolympics.com/>

**TIME magazine** feature story on Bryan Johnson.

<https://time.com/6315607/bryan-johnsons-quest-for-immortality/>

**Men's Health** tested three aging tests, including DunedinPACE.

<https://www.aol.com/lifestyle/fast-am-really-aging-took-191900851.html>

**Youtube: LifeNoggin video: Body secrets**

<https://www.youtube.com/watch?v=kcxdtwYsM>

**Application in geroprotection clinical trials:**

<https://www.newsobserver.com/press-releases/article277627968.html>  
<https://www.youngplasmastudy.com/>  
<https://www.youtube.com/@dianginsbergmd-optimalhealth>

**Nature Biotech:** <https://www.nature.com/articles/d41587-023-00008-6>

**UNUSUAL papers:**

[PDF] [DNA methylation clocks struggle to distinguish inflammaging from healthy aging, but feature rectification improves coherence and enhances detection of inflammaging](#)  
CM Skinner, MJ Conboy, IM Conboy - bioRxiv, 2024

This paper completely describes PACE incorrectly, as trained on people age 53 who had heart attacks and strokes. Hmmmmm.....

[PDF] [Methylation Clocks Do Not Predict Age or Alzheimer's Disease Risk Across Genetically Admixed Individuals](#)

S Cruz-González, E Gu, L Gomez, M Mews, JM Vance... - bioRxiv, 2024

This paper claims that Clocks haven't been studied in blacks and Hispanics and don't work in non-white groups. Hmmmmmm.....