Children Growing Up in Socioeconomically Disadvantaged Families and from Marginalized Racial/Ethnic Groups Tend to Have Epigenetic Profiles Associated with a Faster Pace of Biological Aging

Laurel Raffington, Peter T. Tanksley, Aditi Sabhlok, Liza Vinnik, Travis Mallard, Lucy S. King, Bridget Goosby, Kathryn P. Harden, and Elliot M. Tucker-Drob

INTRODUCTION

The cognitive processes children use to learn, pay attention, remember, and make decisions can be harmed if their environment is stressful or lacks learning opportunities. Children endure stressful environments when they experience such things as limited access to high-quality childcare, educational resources, health care, and nutrition. They also experience stressful environments if they are exposed to family stress, toxic chemicals, and neighborhood threats. Children with social advantages in the United States, particularly those with White identity, or White privilege, are less likely to experience stressful environments because they have benefited from generations of social and legal advantages resulting from classism and racism. Children who identify as Black and of Latin American origin (or Latinx), on the other hand, are more likely to experience stressful environments and more limited learning opportunities.

Scientists can measure environmental effects, including stress, by looking at a child's *epigenetic profile*—a score based on markers on the DNA that turn genes "on" or "off." Epigenetic profiles change as children develop, and can be negatively affected by stressful environments. Previous research with adults has shown that certain epigenetic profiles are associated with negative health outcomes such as increased inflammation (an indicator of stress) and a faster pace of biological aging. Adults who have epigenetic markers of a faster pace of biological aging are more likely to develop health issues such as heart disease, cognitive impairments, as well as earlier death, compared to adults with a slower pace of biological aging.

KEY FINDINGS

- Epigenetic profiles of children from disadvantaged backgrounds looked worse than those of other children. That is, children growing up in more socioeconomically disadvantaged families and neighborhoods and children from marginalized racial/ethnic groups exhibited epigenetic profiles that, in previous studies of adults, were associated with:
 - a faster pace of biological aging (see figure),
 - ▶ higher chronic inflammation, and
 - lower cognitive functioning.
- Children's epigenetic profiles were associated with their performance on tests of cognitive and academic skills, including processing speed, general executive function, perceptual reasoning, verbal comprehension, reading, and math. Children with epigenetic profiles associated with chronic inflammation had worse outcomes.
- These findings from epigenetic profiles support the notion that cognitive function, illness, and death in adulthood are partially driven by molecular processes that begin in childhood.

By analyzing epigenetic profiles in children, scientists can investigate whether associations between epigenetics and negative health outcomes found for adults may begin in childhood. If the associations

INTRODUCTION, CONT.

exist, it is evidence that social inequalities experienced in childhood are carried forward into adulthood in the form of worse health outcomes.

This brief reports on a recent study [1] in which the authors took epigenetic samples from the saliva of young people participating in the Texas Twin Project, a study following twins over time. In the current study, 1,183 participated; they ranged in age from 8 to 19 years, and had an average age of 14. The authors created epigenetic profiles and compared them to profiles that were originally developed for adults to predict the pace of biological aging, chronic inflammation, and cognitive function.

Children from disadvantaged backgrounds tended to have epigenetic profiles associated with a faster pace of biological aging, which in turn can lead to worse health and earlier death



POLICY IMPLICATIONS

Epigenetic profiles are a promising tool and can help us link adult aging to early child experiences to better understand how social inequalities become embedded in the body and impact the mind across the lifespan. To decrease disparities in the cognitive and physical health of adults, interventions need to start in childhood. These interventions would seek to improve children's educational opportunities and nutrition, while also seeking to reduce their family stress and exposure to air pollution and other environmental toxicants.

REFERENCE

[1] 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. *Psychological Science*, *34*(2):170-185. https://doi.org/10.1177/09567976221122760

SUGGESTED CITATION

Raffington, L., Tanksley, P.T., Sabhlok, A., Vinnik, L., Mallard, T., King, L.S., Goosby, B., Harden, K.P., & Tucker-Drob, E.M. (2023). Children growing up in socioeconomically disadvantaged families and from marginalized racial/ethnic groups tend to have epigenetic profiles associated with a faster pace of biological aging. *CAPS Research Brief* 2(1). <u>http://dx.doi.org/10.26153/tsw/44489</u>

ABOUT THE AUTHORS

Laurel Raffington, raffington@mpib-berlin.mpg.de, was a postdoctoral scholar at the Population Research Center (2019-2022), The University of Texas at Austin, and is now research group leader of the Max Planck Research Group Biosocial, Max Planck Institute for Human Development; Peter T. Tanksley is a PRC postdoctoral scholar; Aditi Sabhlok is a PhD candidate in the Department of Psychology, UT Austin; Liza Vinnik is a data scientist with the Texas Twin Project, UT Austin; Travis Mallard is a postdoctoral researcher in the Psychiatric & Neurodevelopmental Genetics Unit at Massachusetts General Hospital and Harvard Medical School; Lucy S. King is a postdoctoral fellow at UT Austin and Tulane University School of Medicine; Bridget Goosby is a professor of sociology, Center on Aging and Population Sciences (CAPS) faculty affiliate and PRC graduate training director; Kathryn P. Harden is a professor of psychology and PRC faculty scholar; and Elliot M. Tucker-Drob is a professor of psychology, CAPS faculty affiliate and PRC faculty scholar. Dr.s Harden and Tucker-Drob contributed equally to this study.

ACKNOWLEDGEMENTS

This research was supported by National Institutes of Health (NIH) Grants R01HD083613 and R01HD092548. L. Raffington was supported by the German Research Foundation. B. Goosby, K. P. Harden, and E. M. Tucker-Drob are faculty scholars of the Population Research Center at The University of Texas at Austin, which was supported by NIH Grant P2CHD042849 (Eunice Kennedy Shriver National Institute of Child Health and Human Development). B. Goosby and E. M. Tucker-Drob are members of the Center on Aging and Population Sciences at The University of Texas at Austin, which was supported by NIH Grant P30AG066614 (National Institute of Aging). K. P. Harden and E. M. Tucker-Drob were also supported by Jacobs Foundation Research Fellowships. L. S. King was supported by NIH Fellowship F32HD105285-01.

This brief is published in partnership with UT Austin's Population Research Center, which provides CAPS with high-quality services and resources to facilitate large-scale, population-based aging research.





The overarching mission of The Center on Aging and Population Sciences (CAPS) is to galvanize novel research that illuminates how biological, psychosocial, and environmental factors intersect and cascade throughout the life course to generate disparities in health and well-being at older ages. The Center promotes collaborations among scholars, supports pilot projects to address complex aging and population health issues, and works to grow the number and diversity of researchers who study aging at all career stages. www.liberalarts.utexas.edu/caps

