

EDITORIAL OPEN



Interplay between genes and social environment: from epigenetics to precision medicine

© The Author(s) 2025

Cell Death Discovery (2025)11:293; <https://doi.org/10.1038/s41420-025-02580-z>

Natural and prolonged exposure to environmental factors substantially influences human physiology throughout life at the molecular and functional level. Gene expression, signal transduction, and cellular processes are altered by exposure, while influencing onset of several diseases such as obesity, cancer, and autoimmune disorders. In biomedicine, deconvoluting the relationship between genetic predispositions and environmental factors (GxE) remains a significant challenge, as genetics alone cannot predict an individual's biological response to a specific environmental exposure. Cells, organs, and tissues can retain a "memory" of previous exposures, which, when combined with developmental and aging factors, create an epigenetic pattern that influences future responses [1]. Therefore, GxE interplay cannot be solely unravelled by examining genetics alone, a detailed understanding of cellular status can only be provided by considering epigenetic patterns. Based on this, we recently proposed the concept of an "epigenetic score metre," a tool that can disentangle the relationship between genetic and environmental influences [2].

Social environment could also participate in the GxE. Major support to this postulation comes from evidence that early-life stress represents an important risk factor for physical disorders [3] as well as mental health conditions and cognitive disabilities [4]. Genetic predispositions, when combined with extrinsic influences, even from social environment, can significantly impact an individual's mental health trajectory [5]. Here, we will provide a summary, supported by historical context and a few brief examples, to highlight how social interaction contributes within GxE interplay to psychological and mild psychiatric thought epigenetic mechanisms (Fig. 1). We will also discuss how deconstructing this interplay could have a significant impact on both society and the economy.

EPIGENETICS AND MENTAL HEALTH

Epigenetics refers to the study of how environmental factors can alter gene expression without changing the underlying DNA sequence. These changes can, in turn, influence behaviour, cognition, and personality. The Diathesis/Stress model, introduced by Zubin and Spring in 1977 [6], offers a framework for understanding mental health disorders. According to this model, such conditions arise from the interaction between genetic vulnerabilities (diatheses) and environmental stressors. It suggests that individuals with a genetic predisposition to mental illness may require specific environmental triggers, such as trauma or adversity, to manifest the disorder. This model emphasizes that mental health disorders result not from genetics or environmental factors alone, but from the interplay between the two.

A twin study by Kendler and colleagues [7] highlighted that 37% of the variance in susceptibility to major depression could be attributed to genetic factors, with substantial overlap between the genetic factors influencing depression and other psychiatric disorders. Similarly, a meta-analysis by Hettema and colleagues [8] showed that between 30% and 50% of the variance in anxiety disorders could be explained by genetic predisposition, with variations depending on the specific disorder. These findings underscore the significant role of genetics in mental health. However, environmental factors also contribute to mental health conditions. A more recent adoption study revealed a significant resemblance in the prevalence of major depressive disorder between children and their adoptive families, indicating environmental transmission [7]. Even if children are not genetically related to their adoptive parents, being raised in a household where caregivers, such as step-parents, struggle with depression increases the likelihood of the child developing depression themselves. This supports the idea that environmental factors, such as the mental health of caregivers, can shape an individual's vulnerability to mental illness.

Changes in gene expression within limbic brain have been associated to aberrant epigenetic regulations with potential causative roles in development of depression and stress-related disorders, such as post-traumatic stress disorder and various anxiety disorders [9]. Consistently, antidepressant medications may exert, at least in part, enduring therapeutic effects by being mediated through epigenetic mechanisms [9].

The environmental effects were also shown to be transmissible with mechanisms of epigenetic inheritance. In mice, chronic exposure to psychosocial stress led to a significant decrease in 5-methylcytosine levels in germ cells. Specifically, psychosocial stress appeared to alter DNA methylation in gene regulatory regions that are involved in transcriptional regulation. These findings suggest that chronic stress disrupts the male reproductive system by inducing abnormal epigenetic changes in male germ cells [10]. This indicates how chronic stress negatively affects germ cell development in the testis and that these effects can be inherited by offspring.

Behavioural traits in mammals have been linked to non-genetic transgenerational inheritance involving the germ line, with epigenetic modifications and non-coding RNAs in germ cells playing a key role [11]. These mechanisms appear similar to non-genetic transgenerational inheritance observed in parental exposure to toxicants or metabolic stress, whose offsprings display higher risk of deficits [12–14].

Consistent with epigenetic research in this area [15], development of common mental health disorders appears to be the integration of multiple influences by an interaction between genetic predispositions and environmental factors, including socioeconomic status, prior trauma, and the mental health of caregivers.

Received: 1 April 2025 Revised: 21 May 2025 Accepted: 13 June 2025
Published online: 01 July 2025

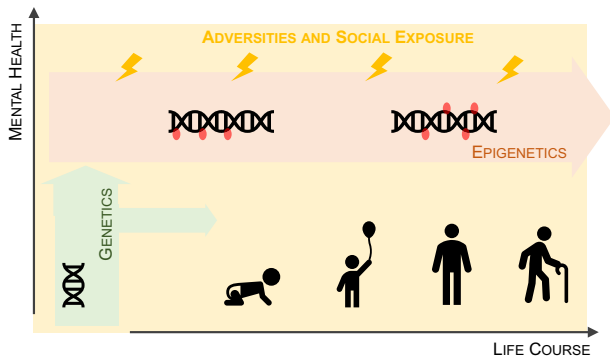


Fig. 1 Schematic representation of the combined influence of genetic and environmental factors on mental health burden. Throughout the life course, various physical and social environmental stressors interact with individual epigenetic mechanisms. Genetic variation further modulates these responses, contributing to diverse mental health outcomes.

HEALTHCARE BURDEN OF MENTAL HEALTH CONDITIONS

The societal impact of mental health disorders, both common and severe, is profound. Research has shown that mental health problems, including common conditions like depression and anxiety, and more severe conditions such as psychosis, place a significant burden on healthcare systems [16, 17]. This burden arises from high hospitalization rates, comorbidity with physical long-term conditions, and the need for prolonged psychological and pharmacological interventions. In 2019 alone, major depression accounted for a staggering \$333.7 billion in economic costs in the United States. Moreover, mental health conditions can severely affect an individual's ability to work, contributing to economic loss. Studies have shown that individuals with depression experience an average loss of 27.2 working days per year, further exacerbating the societal and economic costs associated with these conditions [18, 19].

These findings highlight the importance of understanding and predicting the GxE interactions that contribute to mental health disturbances. Such understanding is vital for preventing the spread of these conditions and reducing their impact on society.

The implementation of predictive strategies in public health services, such as predictive mathematical modelling based on epigenetic profiling capable of estimating an individual's likelihood of developing specific mental health conditions, could enable targeted and effective interventions. By screening high-risk populations and offering early interventions, it would be possible to mitigate the onset and severity of mental health issues. Early intervention, particularly for patients experiencing psychosis, has been shown to reduce symptoms and minimize long-term impacts on functioning [20].

PRECISION MEDICINE IN MENTAL HEALTH TREATMENT

Recent advances in psychiatric research have suggested that precision medicine, tailored to an individual's unique genetic and environmental profile, could play a transformative role in mental health care. As seen in other areas of medicine, such as oncology and cardiology, precision medicine has the potential to improve outcomes by providing individualized treatments based on genetic makeup and environmental exposure. In psychiatry, this could involve personalized pharmacological interventions, accounting for both genetic vulnerabilities and environmental exposures, to optimize treatment efficacy.

Incorporating an epigenetic approach into precision medicine could offer new opportunities for mental health care. By identifying high-risk individuals early, and tailoring interventions to their specific needs, healthcare systems can provide more effective treatments, potentially improving the quality of life for

patients while reducing the long-term costs associated with mental health disorders. This approach could also facilitate greater integration of affected individuals into the workforce, reducing absenteeism and improving economic outcomes.

The interaction between genetic and environmental factors plays a crucial role in the development of mental health disturbances. Epigenetics can provide a valuable framework for understanding how environmental influences shape gene expression and, consequently, mental health outcomes. As research continues to unravel these relationships, the application of predictive models and precision medicine in mental health care holds promise for early intervention and more effective treatments. By incorporating epigenetic insights into public health strategies and mental health care, it is possible to improve patient outcomes, reduce healthcare costs, and alleviate the societal burden of mental health disorders.

Sabrina Caporali¹, Simone Russo², Marcel Leist³
Petra H. Wirtz^{4,5} and Ivano Amelio¹✉

¹Chair for Systems Toxicology, University of Konstanz, Konstanz, Germany. ²School of Psychology & Vision Sciences - George Davis Centre, University of Leicester, Leicester, England, UK. ³Department of *in vitro* Toxicology and Biomedicine, University of Konstanz, Konstanz, Germany. ⁴Centre for the Advanced Study of Collective Behaviour, University of Konstanz, Konstanz, Germany. ⁵Biological Work and Health Psychology, Department of Psychology, University of Konstanz, Konstanz, Germany.
✉email: ivano.amelio@uni-konstanz.de

REFERENCES

- Wu H, Eckhardt CM, Baccarelli AA. Molecular mechanisms of environmental exposures and human disease. *Nat Rev Genet.* 2023;24:332–44.
- Butera A, Smirnova L, Ferrando-May E, Hartung T, Brunner T, Leist M, et al. Deconvoluting gene and environment interactions to develop an “epigenetic score meter” of disease. *EMBO Mol Med.* 2023;15:e18208.
- Nelson CA, Scott RD, Bhutta ZA, Harris NB, Danese A, Samara M. Adversity in childhood is linked to mental and physical health throughout life. *BMJ.* 2020;371:m3048.
- Birnie MT, Baram TZ. The evolving neurobiology of early-life stress. *Neuron* 2025;113:1474–90.
- Klengel T, Binder EB. Epigenetics of stress-related psychiatric disorders and gene x environment interactions. *Neuron.* 2015;86:1343–57.
- Zubin J, Spring B. Vulnerability—a new view of schizophrenia. *J Abnorm Psychol.* 1977;86:103–26.
- Kendler KS, Ohlsson H, Sundquist K, Sundquist J. Sources of parent-offspring resemblance for major depression in a national swedish extended adoption study. *JAMA Psychiatry.* 2018;75:194–200.
- Hettema JM, Neale MC, Kendler KS. A review and meta-analysis of the genetic epidemiology of anxiety disorders. *Am J Psychiatry.* 2001;158:1568–78.
- Vialou V, Feng J, Robison AJ, Nestler EJ. Epigenetic mechanisms of depression and antidepressant action. *Annu Rev Pharmacol Toxicol.* 2013;53:59–87.
- Ohno H, Yamamuro Y, Aizawa S. Chronic social defeat stress alters DNA methylation profiles of male germ cells in mice. *Biochem Biophys Res Commun.* 2025;758:151650.
- Bohacek J, Mansuy IM. Molecular insights into transgenerational non-genetic inheritance of acquired behaviours. *Nat Rev Genet.* 2015;16:641–52.
- Hua W, Han X, Li F, Lu L, Sun Y, Hassanian-Moghaddam H, et al. Transgenerational effects of arsenic exposure on learning and memory in rats: crosstalk between arsenic methylation, hippocampal metabolism, and histone modifications. *Environ Sci Technol.* 2024;58:6475–86.
- McCarthy DM, Morgan TJ Jr., Lowe SE, Williamson MJ, Spencer TJ, Biederman J, et al. Nicotine exposure of male mice produces behavioral impairment in multiple generations of descendants. *PLoS Biol.* 2018;16:e2006497.
- Gonzalez-Rodriguez P, Fullgrabe J, Joseph B. The hunger strikes back: an epigenetic memory for autophagy. *Cell Death Differ.* 2023;30:1404–15.
- Franic S, Middeldorp CM, Dolan CV, Ligthart L, Boomsma DI. Childhood and adolescent anxiety and depression: beyond heritability. *J Am Acad Child Adolesc Psychiatry.* 2010;49:820–9.
- Chong HY, Teoh SL, Wu DB, Kotirum S, Chiou CF, Chaiyakunapruk N. Global economic burden of schizophrenia: a systematic review. *Neuropsychiatr Dis Treat.* 2016;12:357–73.

17. Greenberg P, Chitnis A, Louie D, Suthoff E, Chen SY, Maitland J, et al. The economic burden of adults with major depressive disorder in the United States (2019). *Adv Ther.* 2023;40:4460–79.
18. Kessler RC, Akiskal HS, Ames M, Birnbaum H, Greenberg P, Hirschfeld RM, et al. Prevalence and effects of mood disorders on work performance in a nationally representative sample of U.S. workers. *Am J Psychiatry.* 2006;163:1561–8.
19. Marwaha S, Johnson S. Schizophrenia and employment - a review. *Soc Psychiatry Psychiatr Epidemiol.* 2004;39:337–49.
20. McGorry PD, Killackey E, Yung A. Early intervention in psychosis: concepts, evidence and future directions. *World Psychiatry.* 2008;7:148–56.

ACKNOWLEDGEMENTS

The authors would like to thank all members of the Amelio, Leist, and Wirtz groups for their valuable discussions.

AUTHOR CONTRIBUTIONS

S. Caporali: Writing - original draft; S. Russo: Writing - original draft; M. Leist: Writing - review & editing; I. Amelio: Supervision, writing - review & editing, funding acquisition; P.H. Wirtz: Supervision, writing - review & editing, funding acquisition.

FUNDING

This work has been supported by the DFG to IA, (under the TRR353 “Death Decision” projects A05), the Carl Zeiss Stiftung to IA (Endowed Professorship, #15972218, 2022–2027; Prisma Program, #P2022-5-003), by the cooperation between Carl Zeiss Stiftung and German Scholars Organization with the Fund for international researchers to IA (#15978021) and by the MWK funding to IA. (Az. MWK31-7532-522/1/6).

COMPETING INTERESTS

I. Amelio is Editor-in-Chief of *Cell Death Discovery*. The authors declare no other competing interests.

ADDITIONAL INFORMATION

Correspondence and requests for materials should be addressed to Ivano Amelio.

Reprints and permission information is available at <http://www.nature.com/reprints>

Publisher's note Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.



Open Access This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons licence, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons licence, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons licence and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this licence, visit <http://creativecommons.org/licenses/by/4.0/>.

© The Author(s) 2025