

OPEN ACCESS

EDITED AND REVIEWED BY Raul Mostoslavsky, Massachusetts General Hospital Cancer Center, United States

*CORRESPONDENCE Rasime Kalkan, ⋈ kalkanr@yahoo.com

RECEIVED 31 October 2025 REVISED 05 November 2025 ACCEPTED 10 November 2025 PUBLISHED 18 November 2025

CITATION

Kalkan R, Reinach PS, Harvanek Z and Yan D (2025) Editorial: Epigenetic basis of circadian rhythm and metabolism.

Front. Epigenet. Epigenom. 3:1736716.

doi: 10.3389/freae.2025.1736716

COPYRIGHT

© 2025 Kalkan, Reinach, Harvanek and Yan. This is an open-access article distributed under the terms of the Creative Commons Attribution License (CC BY). The use, distribution or reproduction in other forums is permitted, provided the original author(s) and the copyright owner(s) are credited and that the original publication in this journal is cited, in accordance with accepted academic practice. No use, distribution or reproduction is permitted which does not comply with these terms.

Editorial: Epigenetic basis of circadian rhythm and metabolism

Rasime Kalkan¹*, Peter S. Reinach², Zachary Harvanek³ and Dongsheng Yan⁴

¹Faculty of Medicine, European University of Lefke, Lefke, Türkiye, ²Basic Medical College, Wenzhou Medical University, Wenzhou, China, ³Yale University, New Haven, United States, ⁴Wenzhou Medical University, Wenzhou, China

KEYWORDS

circadian rhythm, metabolism, hormones, sleep, regulation

Editorial on the Research Topic

Editorial: Epigenetic basis of circadian rhythm and metabolism

Circadian rhythms play an important role in regulating several physiological and metabolic processes. The internal molecular clock plays a pivotal role in maintaining including sleep–wake cycles, hormone secretion, body temperature, and metabolism rhythms. Circadian rhythms are changes in the body, brain, and behavior that occur on a 24-h cycle. Circadian rhythms affect several vital bodily processes, including hormone production, digestion, eating patterns, and body temperature regulation (Takahashi, 2017). The clock machinery has a complex transcriptional–translational feedback loop and clock genes like CLOCK, BMAL1, PER, and CRY, together regulate the rhythmic expression of thousands of downstream genes, many of which participate in metabolic pathways.

Environmental factors, including diet, shift work, and ageing, affect epigenetic reprogramming of the circadian clock. Growing evidence indicates that pharmacological manipulation of chromatin modifiers or sirtuins can partially restore circadian and metabolic balance. Thus, therapeutic interventions against metabolic and age-related diseases might be achieved by targeting the epigenetic-circadian-metabolic network. The goal of this Research Topic is to compile the most recent discoveries and developments in chronobiology and epigenetics.

There is an interaction between epigenetic modifications and circadian rhythms (Aguilar-Arnal and Sassone-Corsi, 2013; Dhaka et al., 2025). Mutations or epigenetic alterations on core circadian clock genes (BMAL1, CLOCK, CRY, PER) can cause alterations in the regulation of the brain, heart, muscle, liver, adipose tissue, pancreas, intestine, and, immune system, glucose homeostasis, lipid metabolism, the risk of hospitalized respiratory infections, and myocardial infarction (Guo et al.; Zhu et al.; Reinke and Asher, 2019; Panda, 2016).

Christopher et al. demonstrated that histone deacetylase 3 (HDAC3) regulates the diurnal rhythms of claudin expression and intestinal permeability. The study showed that HDAC3 is an epigenetic regulator that represses claudin expression by deacetylating histones, thereby influencing diurnal rhythms and intestinal permeability. Claudin genes exhibit diurnal rhythmicity, with expression levels fluctuating across the daynight cycle, impacting intestinal permeability. HDAC3 represses claudin expression and promotes diurnal rhythms by deacetylating histones, particularly H3K9ac and H3K27ac(Christopher et al.). Loss of HDAC3 in IECs increases claudin expression,

Kalkan et al. 10.3389/freae.2025.1736716

disrupts diurnal rhythms, and leads to higher intestinal permeability. This study highlights the importance of circadian regulation in maintaining intestinal homeostasis and suggests that targeting HDAC3 could offer therapeutic potential for gastrointestinal disorders linked to circadian disruption (Christopher et al.).

Zhu and colleagues showed that sleep behaviours, including sleep duration, insomnia, chronotype, and daytime sleepiness, were also linked to respiratory infection risk, with healthier sleep behaviours associated with lower risks. Chronotype significantly modified the relationship between Hb and respiratory infection risk, with late chronotypes showing a stronger association between abnormal Hb levels and respiratory infection risk. Their study highlights the importance of maintaining balanced Hb levels and healthy sleep behaviors to reduce the risk of respiratory infections. However, it notes limitations such as reliance on self-reported sleep data, single Hb measurements, and the focus on hospitalized cases rather than community infections (Zhu et al.).

In this Research Topic, Gu and colleagues investigated the relationship between Circadian Syndrome (CircS) and Metabolic Syndrome (MetS) with psoriasis, comparing their predictive abilities using data from the NHANES surveys (2005–2006 and 2009–2014). They showed CircS provides a more comprehensive framework for understanding psoriasis risk compared to MetS, emphasizing the importance of circadian rhythm disruptions and lifestyle factors in psoriasis development. The study highlights the need to monitor CircS components—particularly elevated blood pressure, depressive symptoms, and elevated waist circumference—to better manage psoriasis risk (Gu et al.).

This Research Topic brings together studies investigating the epigenetic regulation of circadian rhythms and metabolism in a variety of biological systems. We seek to discuss new mechanisms, molecular interactions, and translational insights that continue to advance our understanding of how the epigenome regulates temporal and metabolic physiology. Together, these contributions will advance the time-metabolism relationship and open new avenues for chronotherapy and epigenetic-based interventions in metabolic disease. This Research Topic is intended to provide context and provoke further work in the field of epigenetics and circadian rhythm.

References

Aguilar-Arnal, L., and Sassone-Corsi, P. (2013). Chromatin landscape and circadian dynamics: spatial and temporal organization of clock transcription. *Proc. Natl. Acad. Sci. U S A.* 112 (22), 6863–6870. doi:10.1073/pnas.1411264111

Dhaka, P., Neha, Kumar, P., Hossain, C. M., and Parvez, S. (2025). The interplay between circadian rhythms and aging: molecular mechanisms and therapeutic strategies. *Biogerontology* 26 (5), 173. doi:10.1007/s10522-025-10301-3

Author contributions

RK: Writing – review and editing, Writing – original draft. PR: Writing – review and editing, Writing – original draft. ZH: Writing – original draft, Writing – review and editing. DY: Writing – original draft, Writing – review and editing.

Funding

The authors declare that no financial support was received for the research and/or publication of this article.

Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

Generative Al statement

The authors declare that no Generative AI was used in the creation of this manuscript.

Any alternative text (alt text) provided alongside figures in this article has been generated by Frontiers with the support of artificial intelligence and reasonable efforts have been made to ensure accuracy, including review by the authors wherever possible. If you identify any issues, please contact us.

Publisher's note

All claims expressed in this article are solely those of the authors and do not necessarily represent those of their affiliated organizations, or those of the publisher, the editors and the reviewers. Any product that may be evaluated in this article, or claim that may be made by its manufacturer, is not guaranteed or endorsed by the publisher.

Panda, S. (2016). Circadian physiology of metabolism. *Science* 354, 1008–1015. doi:10. 1126/science.aah4967

Reinke, H., and Asher, G. (2019). Crosstalk between metabolism and circadian clocks. *Nat. Rev. Mol. Cell Biol.* 20, 227–241. doi:10.1038/s41580-018-0096-9

Takahashi, J. S. (2017). Transcriptional architecture of the Mammalian circadian clock. *Nat. Rev. Genet.* 18, 164–179. doi:10.1038/nrg.2016.150