

The Exposome journal on hiatus: not even a flesh wound

Gary W. Miller, PhD^{1,*} 

¹Department of Environmental Health Sciences, Mailman School of Public Health, Columbia University, New York, NY, United States

*To whom correspondence should be addressed: Email: gary.miller@columbia.edu

In a memorable scene in Monty Python's *Search of the Holy Grail*, King Arthur is challenged by the Black Knight. After a brief battle, King Arthur severs both of the knight's arms. The knight dismisses the damage as "merely a flesh wound" and continues fighting on. It is hard to defend the knight's detachment from reality, but one must admire his spirit. The ending of the Oxford University Press era of the *Exposome* journal is a slight setback, but I would argue that it isn't even a flesh wound. Many journals have gone on hiatus while new publishing agreements are established. Others have completely ceased operations because the field is undergoing a contraction. But the *exposome* and *exposomics* are not on a downward trajectory. Quite the opposite. The past year has been one of the most exciting for the field, and I AM confident that we will have a new publisher soon.

The ever-changing publishing landscape has put significant pressure on publishers. As the editor, I would have preferred to see greater visibility and more submissions. One can argue about the validity and utility of impact factors, but the vagaries of the bodies that oversee impact factors have some peculiar rules that did not work to our advantage. For example, the assessment criteria determining whether a journal can be awarded an Impact Factor limits the number of papers that include associate editors and editorial board members as co-authors. This is important to identify nefarious paper mills who are cheating the system, but inappropriate for a reputable journal that included many of the most active investigators in the growing field among their associate editors and editorial board. As our field grew, we were unable to attract a sufficient number of submissions to earn an impact factor and meet the publisher's financial targets.

The field of *exposomics* is witnessing exponential growth. Support from numerous funding agencies is swelling, *exposome* technologies are booming, and our anxiety over how to interpret the potential zettabytes of data is being allayed by the therapeutic salve of artificial intelligence. In the U.S. the National Institutes of Health is increasing their support through a variety of grant mechanisms. In the European Union there is strong support for infrastructure and research projects. In the past year there have been multiple high-profile *exposome* papers published in a variety of well-established journals.¹⁻⁹ The *exposome* has also been in the news with significant coverage from *The Economist*, *WIRED*, *the Financial Times*, *the American Association of Medical Colleges*, *the Proceedings of the National Academy of Science*, *the Journal of the American Medical Association*, *Undark*, and many more.

The editorial team at *Exposome* set a high bar on scientific quality and the papers we have published have been and will continue to be influential. Indeed, there have been many exceptional papers published in the journal,¹⁰⁻¹³ including the 20th anniversary paper by Christopher Wild.¹⁴ Some of these papers documented novel methods for the field^{15,16} while others demonstrated the utility of existing methods to explain how *exposomics* can lend new insight to human health concerns.¹⁷⁻¹⁹ The opening and other editorials were highly read.²⁰⁻²² The *Exposome* was also the outlet for several papers that resulted from the National Institute of Environmental Health Sciences Catalytic Workshop Series on the *Exposome*.^{12,13,23-26} The series was attended by over 400 people and dozens of participants helped craft several papers that appeared in the *Journal*.

I have drawn upon Thomas Kuhn's work *The Structure of Scientific Revolutions* numerous times over my career. Science does not follow a straight line. It is punctuated by disruptions and discoveries. In between, there are often periods of steady progress, which he refers to as "normal science." However, as the existing approaches to the scientific problems are challenged, these periods of normality are interrupted by what, at times, can feel like crisis or chaos. I have argued that *exposomics* for the past few years has been experiencing the crisis mode Kuhn described. As the field develops its scientific constructs and principles, there is considerable uncertainty and confusion. Rather than being bad, this phase portends exciting advancements as new fields find their way. Innovation do not always neatly fit into pre-existing boxes.

I AM confident that *Exposome* will be back with a new publishing home soon and we are actively engaged with several publishers. I want to thank Oxford University Press for taking a chance on the journal under its new open access platform and for their commitment to make sure that all of the content of the *Exposome* journal is available in perpetuity. I view the end of the *Exposome* journal era with Oxford University Press not as a flesh wound, but more like a minor bruise. To paraphrase another line from Monty Python, this time from *The Meaning of Life*, "I'm not dead yet ... I'm getting better." The field is stronger than when we started, we weathered some major changes in the publication landscape, and have provided an outlet for dozens of high-quality manuscripts for the field. Now, we must find the most appropriate publication framework for the journal. I remain steadfast that a standalone journal is necessary for the field and

Received: March 8, 2026; Accepted: March 9, 2026

© The Author(s) 2026. Published by Oxford University Press.

This is an Open Access article distributed under the terms of the Creative Commons Attribution-NonCommercial License (<https://creativecommons.org/licenses/by-nc/4.0/>), which permits non-commercial re-use, distribution, and reproduction in any medium, provided the original work is properly cited. For commercial re-use, please contact reprints@oup.com for reprints and translation rights for reprints. All other permissions can be obtained through our RightsLink service via the Permissions link on the article page on our site—for further information please contact journals.permissions@oup.com.

I will continue to work to find a nurturing home to help the journal and the field flourish.

Acknowledgments

The author would like to thank the Associate Editors, Editorial Board Members, authors, and reviewers who had contributed to the journal, and to the Associate Editors who provided comments and suggestions on this editorial.

Author contributions

Gary W. Miller (Conceptualization [equal], Writing—original draft [equal], Writing—review & editing [equal])

Funding

G.W.M. is supported by National Institutes of Health grant ES036819.

Conflicts of interest

None declared.

References

- Argentieri MA, Amin N, Nevado-Holgado AJ, et al. Integrating the environmental and genetic architectures of aging and mortality. *Nat Med*. 2025;31:1016-1025. <https://doi.org/10.1038/s41591-4-03483-9>.
- Hartung T. How AI can deliver the human exposome project. *Nat Med*. 2025;31:1738-1738. <https://doi.org/10.1038/s41591-5-03749-w>.
- Chung MK, Rappaport SM, Wheelock CE, et al. Utilizing a biology-driven approach to map the exposome in health and disease: an essential investment to drive the next generation of environmental discovery. *Environ Health Perspect*. 2021;129:85001. <https://doi.org/10.1289/EHP8327>.
- Wishart D, Arndt D, Pon A, et al. T3DB: the toxic exposome database. *Nucleic Acids Res*. 2015;43:D928-D934. <https://doi.org/10.1093/nar/gku1004>.
- Wan M, Simonin EM, Johnson MM, et al. Exposomics: a review of methodologies, applications, and future directions in molecular medicine. *EMBO Mol Med*. 2025;17:599-608. <https://doi.org/10.1038/s44321-5-00191-w>.
- Sarigiannis D, Karakitsios S, Anesti O, et al. Advancing translational exposomics: bridging genome, exposome and personalized medicine. *Hum Genomics*. 2025;19:48. <https://doi.org/10.1186/s40246-5-00761-6>.
- Maitre L, Bustamante M, Hernández-Ferrer C, et al. Multi-omics signatures of the human early life exposome. *Nat Commun*. 2022;13:7024. <https://doi.org/10.1038/s41467-2-34422-2>.
- Jiang C, Wang X, Li X, et al. Dynamic human environmental exposure revealed by longitudinal personal monitoring. *Cell* 2018; 175:277-291.e31. <https://doi.org/10.1016/j.cell.2018.08.060>.
- Gu Y, Feuerstein ML, Lloyd DT, Patel CJ, Johnson CH, Warth B. Quantitative exposomics targeting over 200 toxicants and key biomarkers at the picomolar level. *Environ Sci Technol*. 2025;59:21818-21829. <https://doi.org/10.1021/acs.est.5c04458>
- Nwanaji-Enwerem JC, Khodasevich D, Gladish N, et al. The environmental chemical exposome and health insurance: examining associations and effect modification of epigenetic aging in a representative sample of United States adults. *Exposome* 2026;6:osaf015. <https://doi.org/10.1093/exposome/osaf015>.
- Safarlou CW, Jongsma KR, Vermeulen R, Bredenoord AL. The ethical aspects of exposome research: a systematic review. *Exposome* 2023;3:osad004. <https://doi.org/10.1093/exposome/osad004>.
- Schmitt CP, Stingone JA, Rajasekar A, et al. A roadmap to advance exposomics through federation of data. *Exposome* 2023;3:osad010. <https://doi.org/10.1093/exposome/osad010>.
- Stingone JA, Geller AM, Hood DB, et al.; Members of the Exposomics Consortium. Community-level exposomics: a population-centered approach to address public health concerns. *Exposome* 2023;3:osad009. <https://doi.org/10.1093/exposome/osad009>.
- Wild CP. The exposome at 20: a personal account. *Exposome*. 2025;5:osaf003. <https://doi.org/doi.org/10.1093/exposome/osaf003>
- Koelmel JP, Stelben P, Godri D, et al. Interactive software for visualization of nontargeted mass spectrometry data—FluoroMatch visualizer. *Exposome*. 2022;2:osac006. <https://doi.org/10.1093/exposome/osac006>.
- Koelmel JP, Xie H, Price EJ, et al. An actionable annotation scoring framework for gas chromatography-high-resolution mass spectrometry. *Exposome* 2022;2:osac007. <https://doi.org/10.1093/exposome/osac007>.
- Haddad N, Lequy E, Zins M, Goldberg M, Makris KC. Methodological advances in designing a longitudinal urban exposome study: the UrbanX exposome study for the city of Paris within the CONSTANCES cohort. *Exposome*. 2026;6:osaf019. <https://doi.org/10.1093/exposome/osaf019>.
- Pfundt K, Hernandez VV, Warth B. Dried milk spots: a viable approach for assessing the chemical exposome in mothers and their infants by targeted LC-MS/MS. *Exposome*. 2026;6:osag001. <https://doi.org/doi.org/10.1093/exposome/osag001>
- Singh RR, Duarte-Hospital C, Mizuno Y, et al. Analytical and cheminformatic framework for studying drugs and their metabolites in human plasma using high resolution mass spectrometry. *Exposome*. 2026;6:osag004. <https://doi.org/doi.org/10.1093/exposome/osag004>
- Miller GW. Exposome: a new field, a new journal. *Exposome* 2021;1:osab001. <https://doi.org/10.1093/exposome/osab001>.
- Miller GW. Integrating the exposome into a multi-omic research framework. *Exposome* 2021;1:osab002. <https://doi.org/10.1093/exposome/osab002>.
- Miller GW. Exposomics: perfection not required. *Exposome* 2024; 4:osae006. <https://doi.org/10.1093/exposome/osae006>.
- Wright RO, Makris KC, Natsiavas P, et al.; Members of the Exposomics Consortium. A long and winding road: culture change on data sharing in exposomics. *Exposome*. 2024;4:osae004. <https://doi.org/10.1093/exposome/osae004>.
- Miller GW. The exposome at NIEHS: from workshops to manuscripts. *Exposome* 2023;3:osad011. <https://doi.org/10.1093/exposome/osad011>.
- Chung MK, House JS, Akhtari FS, et al.; Members of the Exposomics Consortium. Decoding the exposome: data science methodologies and implications in exposome-wide association studies (ExWASs). *Exposome* 2024;4:osae001. <https://doi.org/10.1093/exposome/osae001>.
- Rushing BR, Thessen AE, Soliman GA, Ramesh A, Sumner SCJ; Members of the Exposomics Consortium. The exposome and nutritional pharmacology and toxicology: a new application for metabolomics. *Exposome* 2023;3:osad008. <https://doi.org/10.1093/exposome/osad008>.

© The Author(s) 2026. Published by Oxford University Press.

This is an Open Access article distributed under the terms of the Creative Commons Attribution-NonCommercial License (<https://creativecommons.org/licenses/by-nc/4.0/>), which permits non-commercial re-use, distribution, and reproduction in any medium, provided the original work is properly cited. For commercial re-use, please contact reprints@oup.com for reprints and translation rights for reprints. All other permissions can be obtained through our RightsLink service via the Permissions link on the article page on our site—for further information please contact journals.permissions@oup.com.

Exposome, 2026, 6, 1–2

<https://doi.org/10.1093/exposome/osag011>

Editorial