



Editorial

Is accelerated biological aging the hidden link between physical frailty, social deficits, cognitive impairment and risk of incident diseases?



Over the past decades, the interplay between biological aging and age-related diseases and conditions has become a central focus of geriatric and gerontological research. According to the geroscience hypothesis, biological aging is due to specific mechanisms, such as mitochondrial dysfunction, genomic instability, impaired proteostasis, and chronic low-grade inflammation. These processes progressively lead to multisystem dysregulation across metabolic, immune, endocrine, and repair pathways and provide the biological substrate on which age-related diseases ultimately develop [1–3].

In this framework, physical frailty has been largely regarded as the clinical expression of accelerated biological aging, marking a deviation from the trajectory of normal aging [4,5].

In this issue of the Journal of Nutrition, Health and Ageing, Pei et al. [6] aimed to investigate the relationship between physical frailty, social deficits, cognitive impairment and the risk of developing 45 incident non-communicable diseases (NCDs) using the cohort of the UK Biobank. Moreover, the study explores whether accelerated biological aging mediates these associations, applying three validated indices: the Klemmer–Doubal Method, PhenoAge, and Homeostatic Dysregulation.

The authors reported that impairments across each domain (physical, cognitive, and social) were independently associated with a higher risk of multiple NCDs, ranging from cardiovascular and metabolic disorders to neuropsychiatric and musculoskeletal conditions. The risk increased substantially when deficits coexisted across domains, underscoring the cumulative effect of multidimensional vulnerability. These findings are consistent with previous studies showing that the coexistence of frailty components across multiple domains confers a greater risk of adverse outcomes than single-domain deficits alone [7,8].

In the mediation analyses, accelerated biological aging explained up to 50% of the association between impairment in different domains and disease onset.

On this basis, Pei et al. proposed a conceptual model in which physical, social and cognitive impairments accelerate biological aging, thereby amplifying the risk of non-communicable diseases, a pathway summarized by the authors as “frailty → accelerated biological aging → NCDs.”

As Pei and colleagues correctly acknowledge, however, causality cannot be inferred from their data. The association between physical frailty, cognitive impairment, and social deficits with biological age was evaluated only at baseline, within a cross-sectional design, which precludes establishing temporal directionality.

Moreover, although the indices used to estimate biological age have been validated, several of the underlying biomarkers (e.g., blood pressure, glucose, C-reactive protein, BMI) may also be components or

consequences of these impairments themselves.

In particular, for physical frailty and cognitive impairment, current evidence more strongly supports the reverse direction of influence, that is, accelerated biological aging leading to these conditions, rather than these impairments inducing accelerated aging.

Conversely, the pathway proposed by Pei et al. appears particularly plausible for social deficits, where psychosocial stressors and reduced social participation can influence neuroendocrine, inflammatory, and metabolic pathways involved in biological aging [9–11].

From a clinical perspective, these findings highlight the importance of a comprehensive assessment that includes not only physical and cognitive, but also social dimensions. Although Pei et al. referred to social deficits rather than social frailty, as no validated instrument was used, their results highlight the broader relevance that impairments in the social dimension may have in determining health trajectories.

Social frailty has increasingly been recognized as a key determinant of health and functional decline in later life. Its prevalence among community-dwelling older adults is estimated around 20%, higher than that of physical frailty, and it is consistently associated with disability, depression, and mortality [12]. In the English Longitudinal Study of Ageing, Ragusa et al. found that social frailty, particularly poverty and living alone, was independently linked to all-cause mortality, even after adjustment for multimorbidity and depressive symptoms [13]. Similarly, Goto et al. reported that social frailty nearly doubled the risk of mortality and functional disability across longitudinal studies [14].

Emerging evidence further connects social adversity with biological aging. Kivimäki et al. demonstrated that lifelong social disadvantage was associated with proteomic signatures of accelerated immune aging linked to upregulation of NF-κB-related inflammatory pathways [15]. This convergence between social and biological mechanisms reinforces the view that social frailty is not merely psychosocial, but a state with measurable biological correlates.

Addressing social frailty, by promoting social participation, community engagement, and supportive networks, may therefore represent not only a social but also a medical intervention. Integrating validated measures of social frailty into clinical and epidemiological studies will be essential to strengthen the evidence and to support its broader implementation in research and practice.

Finally, in this study, the evidence that impairments in one or more domains are associated with a higher risk of incident non-communicable diseases, was already evident before old age, strongly suggesting that a multidimensional assessment may be appropriate even before advanced age.

<https://doi.org/10.1016/j.jnha.2025.100717>

Received 29 October 2025

Available online 7 November 2025

1279-7707/© 2025 The Authors. Published by Elsevier Masson SAS on behalf of SERDI Publisher. This is an open access article under the CC BY-NC-ND license (<http://creativecommons.org/licenses/by-nc-nd/4.0/>).

Declaration of competing interest

The authors have no conflict of interest to declare.

References

- [1] Kennedy BK, Berger SL, Brunet A, Campisi J, Cuervo AM, Epel ES, et al. Geroscience: linking aging to chronic disease. *Cell*. 2014;159(4):709–13. <https://doi.org/10.1016/j.cell.2014.10.039>.
- [2] Ferrucci L, Gonzalez-Freire M, Fabbri E, Simonsick E, Tanaka T, Moore Z, et al. Measuring biological aging in humans: a quest. *Aging Cell*. 2020;19(3):e13080. <https://doi.org/10.1111/ace1.13080>.
- [3] Kritchevsky SB, Cummings SR. Geroscience: a translational review. *JAMA*. 2025; 334(12):1094–102. <https://doi.org/10.1001/jama.2025.11289>.
- [4] Salvioli S, Basile MS, Bencivenga L, Carrino S, Conte M, Damanti S, et al. Biomarkers of aging in frailty and age-associated disorders: State of the art and future perspective. *Ageing Res Rev*. 2023;91:102044. <https://doi.org/10.1016/j.arr.2023.102044>.
- [5] Kim DH, Rockwood K. Frailty in older adults. *N Engl J Med*. 2024;391(6):538–48. <https://doi.org/10.1056/NEJMra2301292>.
- [6] Pei B, Yu Y, Shen X, Jia Y, Wang J, Zhang Y, et al. Associations of physical frailty, social deficits, and cognitive impairment with risk of 45 incident non-communicable diseases: the mediating role of accelerated biological aging—a prospective cohort study. *J Nutr Health Aging*. 2025;29(11):100672. <https://doi.org/10.1016/j.jnha.2025.100672>.
- [7] Lee Y, Kim E, Yun J, Chuck KW. The influence of multiple frailty profiles on institutionalization and all-cause mortality in community-living older adults. *J Cachexia Sarcopenia Muscle*. 2022;13(5):2322–30. <https://doi.org/10.1002/jcsm.13033>.
- [8] Shimoda T, Tomida K, Nakajima C, Kawakami A, Tsutsumimoto K, Shimada H. Prevalence and prognostic impact of multiple frailty domain in Japanese older adults. *J Am Med Dir Assoc*. 2024;25(11):105238. <https://doi.org/10.1016/j.jamda.2024.105238>. Epub 2024 Sep 3.
- [9] Ong AD, Mann FD, Kubzansky LD. Cumulative social advantage is associated with slower epigenetic aging and lower systemic inflammation. *Brain Behav Immun Health*. 2025;48:101096. <https://doi.org/10.1016/j.bbih.2025.101096>.
- [10] Tam LM, Hocker K, David T, Williams EM. The influence of social dynamics on biological aging and the health of historically marginalized populations: a biopsychosocial model for health disparities. *Int J Environ Res Public Health*. 2024;21(5):554. <https://doi.org/10.3390/ijerph21050554>.
- [11] Hamilton OS, Steptoe A. Socioeconomic determinants of inflammation and neuroendocrine activity: a longitudinal analysis of compositional and contextual effects. *Brain Behav Immun*. 2023;107:276–85. <https://doi.org/10.1016/j.bbi.2022.10.010>.
- [12] Yamada M, Arai H. Understanding social frailty. *Arch Gerontol Geriatr*. 2023 Dec; 115:105123. <https://doi.org/10.1016/j.archger.2023.105123>.
- [13] Ragusa FS, Veronese N, Smith L, Koyanagi A, Dominguez LJ, Barbagallo M. Social frailty increases the risk of all-cause mortality: a Longitudinal analysis of the English Longitudinal Study of Ageing. *Exp Gerontol*. 2022 Oct 1;167:111901. <https://doi.org/10.1016/j.exger.2022.111901>.
- [14] Goto T, Kishimoto T, Fujiwara S, Shirayama Y, Ichikawa T. Social frailty as a predictor of all-cause mortality and functional disability: a systematic review and meta-analysis. *Sci Rep*. 2024 Feb 10;14(1):3410. <https://doi.org/10.1038/s41598-024-53984-3>.
- [15] Kivimäki M, Pentti J, Frank P, Liu F, Blake A, Nyberg ST, et al. Social disadvantage accelerates aging. *Nat Med*. 2025 May;31(5):1635–43. <https://doi.org/10.1038/s41591-025-03563-4>.

Massimiliano Fedecostante^a, Jacopo Sabbatinelli^{b,c},
Antonio Cherubini^{a,b,*,1}

^a *Geriatrics, Accettazione Geriatrica e Centro di Ricerca per l'Invecchiamento, IRCCS INRCA, Via Della Montagnola n. 81, 60127, Ancona, Italy*

^b *Department of Clinical and Molecular Sciences (DISCLIMO), Università Politecnica Delle Marche, Ancona, Italy*

^c *Clinic of Laboratory and Precision Medicine, IRCCS INRCA, 60127 Ancona, Italy*

* Corresponding author.

E-mail address: a.cherubini@inrca.it (A. Cherubini).

¹ Given his role as editorial board member, Antonio Cherubini had no involvement in the peer-review of this article and has no access to information regarding its peer-review. Full responsibility for the editorial process for this article was delegated to another journal editor.