

# Genetic Markers of Biological Age in Multi-component Lifestyle Interventions: Results of a Preliminary Literature Search

M. Gasser<sup>1</sup>, J. Bürgi<sup>2</sup>, W. Korte<sup>2</sup>, M. Hergersberg<sup>2</sup> & T. Kowatsch<sup>1,3,4</sup>

<sup>1</sup>School of Medicine, University of St.Gallen, St.Gallen, Switzerland; <sup>2</sup>Zentrum für Labormedizin St.Gallen, St.Gallen, Switzerland; <sup>3</sup>Institute for Implementation Science in Health Care, University of Zurich, Zurich, Switzerland; <sup>4</sup>Centre for Digital Health Interventions, Department of Management, Technology and Economics, ETH Zurich, Zurich, Switzerland

## Motivation

Mobile health (mHealth) apps can potentially support patients and health care systems (Jakob et al. 2022). Of particular interest is the application of mHealth apps to multi-component lifestyle interventions, such as diet and nutrition, physical activity, sleep and stress management behavior. Such interventions have the potential to slow down or even revert aging processes (Fiorito et al. 2021). Many clinical and laboratory biomarkers have been evaluated to document the aging process and healthy aging (Li et al. 2021). Popular markers are telomere length and epigenetic methylation clocks (Seale et al. 2022). Studies evaluating epigenetic clocks as biomarkers have yielded inconclusive results (Galow & Peleg 2022).

We plan to study the effects of a mHealth intervention on epigenetic clocks as biomarkers of healthy aging. A preliminary literature search was conducted to inform the design of such an intervention.

## Method

A preliminary literature search was conducted that will inform a scoping review. For this purpose, PubMed database was used with the following search terms: lifestyle intervention, epigenetic clock, telomere length and aging biomarker.

## Preliminary Results

## Future Work

As a next step, the scoping review will be finalized. The results will inform the design of a smartphone-based and chatbot-delivered multi-component intervention and the sample size of a randomized controlled trial (RCT). The following data will be collected at the beginning and after the intervention, as well as during a 6-month follow-up of the RCT: vital parameters, serum samples, and stool samples. Epigenetic (biological) age (GrimAge/TruAge) will be determined from saliva and blood cells at the beginning and the end of the intervention, as well as during the follow-up. The metabolome of the blood samples (250 metabolites (NMR von Nightingale (Helsinki); Würtz et al. 2017)) will be determined, too. Adherence to the intervention will be measured continuously via the mHealth app.

## References

- Jakob, R., Harperink, S., Rudolf, A. M., Fleisch, E., Haug, S., Mair, J. L., Salamanca-Sanabria, A., & Kowatsch, T. (2022). Factors Influencing Adherence to mHealth Apps for Prevention or Management of Noncommunicable Diseases: Systematic Review. *J Med Internet Res*, 24(5), e35371. <https://doi.org/10.2196/35371>
- Galow, A.-M., & Peleg, S. (2022). How to Slow down the Ticking Clock: Age-Associated Epigenetic Alterations and Related Interventions to Extend Life Span. *Cells*, 11(3). <https://doi.org/10.3390/cells11030468>
- Seale, K., Horvath, S., Teschendorff, A., Eynon, N., & Voisin, S. (2022). Making sense of the ageing methylome. *Nature Reviews Genetics*, 23(10), 585-605. <https://doi.org/10.1038/s41576-022-00477-6>
- Fiorito, G., Cairni, S., Palli, D., Bendinelli, B., Saieva, C., Ermini, I., Valentini, V., Assedi, M., Rizzolo, P., Ambrogetti, D., Ottini, L., & Masala, G. (2021). DNA methylation-based biomarkers of aging were slowed down in a two-year diet and physical activity intervention trial: the DAMA study. *Aging cell*, 20(10), e13439-e13439. <https://doi.org/10.1111/acel.13439>
- Würtz, P., Kangas, A. J., Soininen, P., Lawlor, D. A., Davey Smith, G., & Ala-Korpela, M. (2017). Quantitative Serum Nuclear Magnetic Resonance Metabolomics in Large-Scale Epidemiology: A Primer on -Omics Technologies. *Am J Epidemiol*, 186(9), 1084-1096. <https://doi.org/10.1093/aje/kwx016>

### Occurrence of Intervention components (IC) in % (published and registered studies N = 39)

Intervention component	N	% of IC overall
Physical activity	13	51%
Diet	13	33%
Supplements	5	15%
Mental health exercises	6	28%
Sleep	3	13%
Combination of ICs	8	28%

### Occurrence of published and registered studies

Type of study	N	% of published / registered studies
Published RTCs	22	56%
Registered RTCs	17	44%

### Biological age biomarkers (published studies only)

Type of epigenetic clock	N	% of age biomarkers	Thereof significant	% signif. per IC
Epigenetic age clocks	6	27%	4	67%
DNAm changes in local cells/tissues	1	5%	1	100%
Telomere length	15	68%	7	47%

### Occurrence of Intervention IC in published studies (N = 17) & significance

Intervention component	N	% of IC published studies	Thereof significant	% signif. per IC
Physical activity	10	45%	6	60%
Diet	11	50%	8	73%
Supplements	4	18%	2	50%
Mental health exercises	6	27%	3	50%
Sleep	3	14%	1	33%
Combination of ICs	6	27%	4	67%

### Details epigenetic age clocks (only published studies)

Epigenetic age clock	N	Occurrence within age clocks
Horvath's DNAmAge clock	3	50%
DNAmGrimAge	1	17%
Hannum's DNAm clock	1	17%
Zbiec-Piekarska's clock	1	17%

### Details epigenetic Telomere length (only published studies)

Type of measurement	N	Occurrence within telomere length
Average Telomere length	15	100%
% change of short telomeres	1	7%

