

Metabolomics-based predictors as markers of health and disease



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Biological Age Biomarkers

To quantify the age-related decline





'Clocks' The rationale for training (omics) age-predictors







Original Article

Circulating Proteomic Signatures of Chronological Age

Cristina Menni,^{1,*} Steven J. Kiddle,^{2,3,*} Massimo Mangino,¹ Ana Viñuela,¹

Attde protection research Measuring Biological Age via Metabonomics: The Metabolic Age Score Johannes Hertel,*^{1,1} Nele Friedrich,^{1,8} Katharina Wittfeld,¹¹ Maik Pietzner,¹ Kathrin Budde,¹

Aim Training and evaluating a Nightingale metabolomics age-predictor



Nightingale Health blood metabolomics

- Highly standardized
- □ Affordable
- 226 metabolites
- Soft cardiovascular focus
- □ Known age-related dynamics



Kirsi Auro^{1,2,3,*}, Anni Joensuu^{1,2,*}, Krista Fischer⁴, Johannes Kettunen^{1,2,5}, Perttu Salo^{1,2}, Hannele Mattsson^{1,2},

Study Population after Quality Control

BBMRI: 25 453 samples from 26 biobanks



Training and evaluating the age-predictor 5 cross-fold validation



predicted age = $\beta_0 + \beta_1 m_1 + \beta_2 m_2 + \dots + \beta_{56} m_{56} + \epsilon$

Training the age-predictor Contributing metabolites



Note: metabolites are scaled and centered before modelling!

Training and evaluating the age-predictor Leave-One-Biobank-Out Cross Validation



Construction of the metaboAge Score The age-independent part of the age-predictor



Evaluating the metaboAge Score Tracking Age-Related Disease



Associations with Clinical Risk Factors

Body Mass Index



 $\Delta age = \beta_0 + \beta_1 BMI + \beta_2$ chronological age + β_3 gender + ϵ

Associations with Current Clinical Endpoints Diabetes Mellitus Status





PROSPER - INCIDENT CARDIOVASCULAR EVENTS

Phenotype	HR	95% CI	<i>p</i> -value
Coronary events	1.25	1.11 - 1.40	2.64 x 10⁻⁰⁴
Cardiovascular events	1.20	1.08 - 1.33	4.86 x 10 ⁻⁰⁴
Vascular mortality	1.57	1.31 - 1.88	8.56 x 10 ⁻⁰⁷
All-cause mortality	1.42	1.25 – 1.61	9.14 x 10 ⁻⁰⁸
Heart failure hospitalisation	1.68	1.37 – 2.06	5.42 x 10 ⁻⁰⁷



Effect independent of BMI, smoking, diabetes and hypertension and medication • Nightingale Health blood metabolomics predicts age reasonably (r = 0.65)

- Comparable with transcriptomics & metabolomics clocks
- Worse than epigenetics clocks (*r* > 0.9). **Does this matter?**
- *MetaboAge* shows promise as an biomarker
 - *MetaboAge* associates significantly with clinical risk factors
 - *MetaboAge* associates significantly with current clinical end points
 - *MetaboAge* associates significantly with future clinical end points
- *MetaboAge* is:
 - Highly standardized
 - Affordable
 - Easy to implement

Discussion **Other clocks**

Small correlation with Horvath's methylation clock





What outcomes do the clocks predict? What do they predict independently?

Small correlation with Deelen's mortality predictor



ARTICLE

https://doi.org/10.1038/s41467-019-11311-9

OPEN A metabolic profile of all-cause mortality risk

identified in an observational study of 44,168 individuals

Joris Deelen tal.#

Discussion Metabolomics as a read-out of intervention success



Other risk factors might be readily predicted And transformed in discrepancy-based predictors





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Application 1: Imputation



 $\Delta Diabetes$, $\Delta Obesity$, ...



Platform	Samples	Measurements
Nighting ale	349	224
Biocrates	356	163
Lipidyzer	307	1107
Metabolon	304	762

Cathelijn Kuijt

Dennis Mook

ELANET (fixed)	(fixed) AGE		BMI		SEX		DIAB	
R-squared	RSQ	df	RSQ	df	RSQ	df	RSQ	df
Nightingale	35,72%	96	36,01%	42	68,91%	58	47,88%	45
Biocrates	47,25%	121	42,17%	96	55,96%	94	55,49%	50
Lipidyzer	51,88%	154	17,51%	104	38,03%	231	35,75%	129
Metabolon	59,50%	162	48,66%	188	77,00%	195	60,23%	135

Some platforms are more predictive for cardio-metabolic variables

Availability Manuscript & tools

metaboAge

Welcome to the webtool for calculating metabolic age (*'metaboAge'*) from raw Nightingale Health 1H-NMR metabolomics data.

Please refer to our manuscript when using metaboAge in your work:

E.B. van den Akker et al. "MetaboAge: a novel biomarker for biological age based on the BBMRI-NL 1H-NMR metabolomics repository" (<u>submitted</u>)



Jurriaan Barkey Wolf

Instructions

For the webtool to work, the input data needs to be provided in a specific format. This format should be similar to the raw data files you received from Nightingale Health.

Please ensure that your dataset looks like this example dataset before submitting.

This means that:



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R-shiny application is coming to compute metaboAge and other scores!

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MIMOmics







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