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Cohort profile: "Biomarkers of Personalized Medicine" (BioPersMed) – A single-centre prospective observational cohort study in Graz/Austria to evaluate novel biomarkers in cardiovascular and metabolic diseases

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Complete List of Authors:	<p>Haudum, Christoph; Center for Biomarker Research in Medicine , ; Medical University of Graz, Department of Internal Medicine</p> <p>Kolesnik, Ewald; Medizinische Universität Graz, Department of Internal Medicine and University Heart Center Graz</p> <p>Colantonio, Caterina ; Medizinische Universität Graz, Department of Internal Medicine and University Heart Center Graz</p> <p>Mursic, Ines; Medical University of Graz, Department of Internal Medicine</p> <p>Url-Michitsch, Marion; Medical University of Graz, Department of Internal Medicine and University Heart Center Graz</p> <p>Tomaschitz, Andreas; Medical University of Graz, Department of Internal Medicine and University Heart Center Graz</p> <p>Glantschnig, Theresa ; Medical University of Graz, Department of Internal Medicine and University Heart Center Graz</p> <p>Hutz, Barbara; Medical University of Graz, Department of Internal Medicine</p> <p>Lind, Alice ; Medical University of Graz, Department of Internal Medicine</p> <p>Schweighofer, Natascha; Medical University of Graz, Department of Internal Medicine</p> <p>Reiter, Clemens ; Medical University of Graz, Department of Internal Medicine and University Heart Center Graz; Medizinische Universität Graz, Department of Radiology</p> <p>Ablasser, Klemens; Medical University of Graz, Department of Internal Medicine and University Heart Center Graz</p> <p>Wallner, Markus; Medical University of Graz, Department of Internal Medicine and University Heart Center Graz; Lewis Katz School of Medicine at Temple University, Cardiovascular Research Center</p> <p>Tripolt, Norbert; Medizinische Universität Graz, Department of Internal Medicine</p> <p>Pieske-Kraigher, Elisabeth; Charité University Medicine, Department of Internal Medicine and Cardiology</p> <p>Madl, Tobias; Medical University of Graz, Gottfried Schatz Research Center; BioTechMed</p> <p>Springer, Alexander; Medical University of Graz, Gottfried Schatz Research Center; BioTechMed</p> <p>Seidel, Gerald; Medical University of Graz, Department of Ophthalmology</p> <p>Wedrich, Andreas; Medizinische Universität Graz, Department of Ophthalmology</p>

	<p>Zirlik, Andreas; Medical University of Graz, Department of Internal Medicine and University Heart Center Graz</p> <p>Krahn, Thomas; Maastricht University, Department of Pharmacology and Personalised Medicine</p> <p>Stauber, Rudolf; Medizinische Universität Graz, Department of Internal Medicine</p> <p>Pieske, Burkert; Charité University Medicine, Department of Internal Medicine and Cardiology,</p> <p>Pieber, Thomas; Medical University of Graz, Department of Internal Medicine; Center for Biomarker Research in Medicine</p> <p>Verheyen, Nicolas; Medizinische Universität Graz, Department of Internal Medicine and University Heart Center Graz</p> <p>Obermayer-Pietsch, Barbara; Medical University of Graz, Department of Internal Medicine</p> <p>Schmidt, Albrecht; Medizinische Universität Graz, Department of Internal Medicine and University Heart Center Graz</p>
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BioPersMed Concept Paper

Cohort profile: “Biomarkers of Personalized Medicine” (BioPersMed) – A single-centre prospective observational cohort study in Graz/Austria to evaluate novel biomarkers in cardiovascular and metabolic diseases

Authors:

Haudum CW^{1,2°}, Kolesnik E^{3°}, Colantonio C³, Mursic I¹, Url-Michitsch M³, Tomaschitz A³, Glantschnig T³, Hutz B¹, Lind A¹, Schweighofer N¹, Reiter C^{3,4}, Ablasser K³, Wallner M^{3,10}, Tripolt N¹, Pieske-Kraigher E⁵, Madl T^{6,11}, Springer A^{6,11}, Seidel G⁷, Wedrich A⁷, Zirlik A³, Krahn T^{8,12}, Stauber R⁹, Pieske B⁵, Pieber TR^{1,2}, Verheyen N³, Obermayer-Pietsch B^{1*§}, Schmidt A^{3*§}

° and * - these authors contributed equally to the manuscript

§ Authors for correspondence and reprint requests

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Affiliations:

¹ Department of Internal Medicine, Division of Endocrinology and Diabetology, Medical University of Graz, Graz, Austria

² Center for Biomarker Research in Medicine (CBmed), Graz, Austria.

³ Department of Internal Medicine and University Heart Center Graz, Division of Cardiology, Medical University of Graz, Graz, Austria

⁴ Department of Radiology, Medical University of Graz, Graz, Austria

⁵ Department of Internal Medicine and Cardiology, Charité University Medicine, Campus Virchow Klinikum and German Heart Center, Berlin, Germany

⁶ Gottfried Schatz Research Center, Chair of Molecular Biology and Biochemistry, Medical University of Graz, Graz, Austria

⁷ Department of Ophthalmology, Medical University of Graz, Graz, Austria.

⁸ Bayer AG, Berlin, Germany

⁹ Department of Internal Medicine, Division of Gastroenterology and Hepatology, Medical University of Graz, Graz, Austria

¹⁰ Lewis Katz School of Medicine, Temple University, Cardiovascular Research Center, Philadelphia, PA, USA

¹¹ BioTechMed-Graz, Graz, Austria.

¹² Department of Pharmacology and Personalised Medicine, Faculty of Health, Medicine and Life Sciences, Maastricht University, Maastricht, Netherlands

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Abstract [244/max 250w]

Purpose

Accumulating evidence points towards a close relationship between cardiovascular, endocrine and metabolic diseases. The BioPersMed Study (Biomarkers of Personalized Medicine) is a single-centre prospective observational cohort study with repetitive examination of participants in 2-year intervals. The aim is to evaluate the predictive impact of various traditional and novel biomarkers of cardiovascular, endocrine and metabolic pathways in asymptomatic individuals at risk for cardiovascular and/or metabolic disease.

Participants:

Between 2010 and 2016, we recruited 1022 regional individuals into the study. Subjects aged 45 years or older presenting with at least one traditional cardiovascular risk factor or manifest type 2 diabetes mellitus were enrolled. The mean age of the participants was 57 ± 8 years, 55 % were female, 18 % had T2DM, 33 % suffered from arterial hypertension, 15 % were smokers, 42 % had hyperlipidaemia, and only 26 % were at low cardiovascular risk according to the Framingham "Systematic COronary Risk Evaluation" (SCORE).

Findings to date:

Study procedures during screening and follow-up visits included a physical examination and comprehensive cardiovascular, endocrine, metabolic, ocular, and laboratory workup with biobanking of blood and urine samples. The variety of assessed biomarkers allows a full phenotyping of individuals at cardiovascular and metabolic risk. Preliminary data from the cohort and relevant biomarker analyses were already used as control population for genomic studies in local and international research cooperations.

Future plans:

Participants will undergo comprehensive cardiovascular, endocrine, and metabolic examinations for the next decades and clinical outcomes will be adjudicated prospectively.

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Strength and Limitations of this study

- The main strength of the BioPersMed cohort is the joint evaluation of cardiovascular, endocrine and metabolic phenotyping including a broad spectrum of highly innovative diagnostic, imaging and functional tools.
- Biobanking with a large number of samples aliquoted and stored at each visit enables a prospective view on candidate biomarkers in the context of a large longitudinal cohort, where specific approaches can be predefined.
- A specifically adapted large electronic data capture system (OpenClinica; www.openclinica.com) and iterated monitoring assures the quality of data entry and delivery as well as the validity and reliability of biomarkers analyses.
- A potential weakness of this study is the wide time range of recruitment due to logistic reasons between 2010 and 2016, with a prolonged follow-up period of study participants to date between 4 and 10 years.
- Some biomarkers are not available for the complete duration of the entire cohort

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Introduction

Cardiovascular (CVD) and metabolic diseases (MD) are globally representing the most important cause of disability and premature death [1]. Next to our genetic programming, modern lifestyle, including the use of tobacco, unhealthy nutritional habits, physical inactivity, and psychosocial stress are major risk factors of CVD and metabolic diseases within different age groups not only promoting excess cardiovascular (CV) and metabolic morbidity but ultimately triggering excess mortality [2,3]. In turn, primary prevention of these diseases has the potential to avoid many of related deaths [4]. However, the initial euphoria about a decline of CVD prevalence at the beginning of this century [5] gradually gives way to a sense of frustrating in the light of increasing numbers of type 2 diabetes mellitus (T2DM) [6], a very high lifetime-risk for the development of heart failure with stable incidence over the last decades [7], a persistent high stroke mortality [8] and the relevant underdiagnosing and/or undertreatment of patients at high risk for CVD related risk factors remaining undiagnosed and un- or undertreated [9]. Therefore, early detection of asymptomatic CV and/or metabolic risk remains a crucial challenge in the prevention of both, onset and progression of CVD as well as of related complications [10].

Considering the multiplicity of risk pathophysiology, an integrative approach is needed to identify novel and to validate established CV and metabolic biomarkers for their scientific and clinical utility. Practical biomarkers are required to facilitate (1) the understanding of underlying mechanisms of disease development, (2) the detection of potential targets for specific preventive therapies, (3) the precise estimation of individual risk and finally (4) the risk-adapted personalized tailoring of therapy. For this purpose, there is an unmet need for a cohort studies recruiting individuals a risk for CVD and/or MD well before clinical manifestation of the diseases.

In the BioPersMed cohort (Biomarkers of Personalized Medicine), we enrolled community dwelling and asymptomatic individuals from the regional communities who were at risk for CVD or MD in order to evaluate the predictive value of various traditional and novel biomarkers. The latter shed light on different pathways of CVD and MD development by use of cutting edge laboratory measurements, advanced imaging techniques, comprehensive genetic investigations, and state of the art functional tests. The BioPersMed cohort is located at the Medical University of Graz (Austria) in a dedicated clinical outpatient research centre and biobank (www.biobank.medunigraz.at). The aim of the study is to evaluate large-scale

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3 screening tools for the improvement of (1) cardiovascular and metabolic risk stratification, (2)
4 early diagnosis of CVD and/or MD (3), individual prediction of clinical outcomes, (4) and long-
5 term monitoring of risk and/or early CV and/or metabolic changes in an apparently healthy
6 but representative at-risk population at high CV and/or metabolic risk in a prospective
7 manner. Ultimately, the data obtained from this cohort aims to facilitate the implementation
8 of risk-adapted personalized interventions in both primary and secondary prevention of CVD
9 and MD.
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Cohort description

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20 The BioPersMed project is designed as a single-centre, prospective, observational study. Only
21 asymptomatic subjects without diagnosed CVD but with at least one traditional CV risk factor
22 were eligible to participate. According to the published European Guidelines on cardiovascular
23 disease prevention in clinical practice, traditional CV risk factors besides of age and gender
24 comprise (1) smoking, (2) elevated total cholesterol levels, and (3) arterial hypertension [11].
25 Moreover, sedentary lifestyle, obesity, social environment, type 1 diabetes mellitus (T1DM)
26 or T2DM, low HDL cholesterol, increased triglyceride levels, elevated fibrinogen,
27 apolipoprotein B (apoB), lipoprotein(a), familial hypercholesterinaemia, increased high
28 sensitivity (hs)CRP, preclinical evidence of atherosclerosis and chronic kidney disease
29 [glomerular filtration rate (eGFR) ≤ 60 mL/min/1.73 m²] were regarded as additional potential
30 CV and MD risk factors. From October 2010 (first patient in) to February 2016 (last patient in),
31 we enrolled a total of 1022 community dwelling adult men and women who live in the greater
32 Graz area via an established recruitment network, consisting of general practitioners,
33 peripheral hospitals, and in most cases through the outpatient clinics of the Departments of
34 Cardiology as well as Endocrinology and Diabetology, respectively.
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47 Subjects presenting with significant non-CVD or who were expected not to be able to
48 complete study specific examinations, were excluded from participation. Ethical approval for
49 the BioPersMed cohort study has been granted by the Ethics Committee of the Medical
50 University of Graz, Austria and is renewed every year (EC Nr. 24-224 ex 11/12). The
51 BioPersMed study was conducted in compliance with Good Clinical Practice Guidelines
52 Procedures (GCP) and complies with the Declaration of Helsinki and the Austrian laws. All
53 participants in the BioPersMed cohort were thoroughly checked for in- and exclusion criteria
54 before the first phenotyping at baseline examination in order to avoid screening failures.
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3 The baseline examinations have been repeated every two years in addition to interim
4 telephone visits, which take place between the on-site visits. A summary of all examinations
5 is shown in figure 1, and a more detailed description can be found in the supplement
6 (Supplemental Tables S1 to S10).
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10 According to the presented scheme, participants will be followed for the next decades and
11 clinical outcomes are adjudicated prospectively. A total of 169 (17 %) participants dropped-
12 out for various reasons. Causes for premature unintended termination of the study ranged
13 from a change in the place of residence (n = 5) to limited personal time or lacking will for
14 continuous study visits (n= 136) or new onset of non-CV related diseases (cancer: 4, accident:
15 3, other: 9). Twelve people have died so far (cancer: 7, sepsis: 3, CVD: 2). In summary, 1022
16 persons were included in the baseline examination and 799 persons attended the first follow-
17 up two years after baseline examination. With September 1st, 2021, 628 persons have
18 completed the second follow-up at four years after baseline visit, 531 persons have completed
19 the third follow-up at six years after the baseline visit, and 225 persons have completed the
20 follow-up of eight years after the baseline visit. A small number of participants skipped one
21 follow-up but decided to continue to participate in the study. This issue explains the
22 discrepancy between the number of drop-outs and the number of missing follow-up visits. A
23 detailed overview of the recruitment process and study protocol is presented in figure 2.
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36 At baseline and at regular two-years-follow-ups, an in-depth diagnostic CV and MD work-up
37 was carried out, laying emphasis on standardization and reproducibility of history taking,
38 questionnaires for health, psychological and sleep issues, physical examination, ECG,
39 laboratory/blood sampling with biobanking, exercise testing (6min walking test, grip strength,
40 spiroergometry), echocardiographic analysis of cardiac structure and function, pulmonary
41 function testing, carotid intima/media-thickness measurement, pulse-wave analysis, and
42 ophthalmologic examinations as well as body composition, bone density including bone and
43 hormonal biomarkers, and oral glucose tolerance testing. The number of examinations
44 performed at each visit increased over time, due to additional new diagnostic tools (e.g., non-
45 mydriatic fundoscopy). A detailed description of all methods used, as well as a concise
46 overview of the assessed data can be found in the supplementary file. Statistics have been
47 calculated using RStudio Version 1.2.5033 (RStudio Inc., United States of America) [12].
48 Normal distribution of data was tested and, if positive, Pearson-correlation calculated. In case
49 of a violation of normal distribution, a non-parametric equivalent was used. A description of
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3 the data is given in the corresponding tables. Before analysis, an 80 % monitoring of the whole
4 data set was performed by certified monitors. Biological and technical outlier were manually
5 identified and corrected. This study conforms to the STROBE-ME recommendations for
6 reporting cohort studies [13].
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Health status

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16 At each visit, a careful assessment of the participants' health status was performed. Data
17 collection documents anthropometric, biochemical, metabolic, hormonal, dietary, physical
18 activity, socioeconomic, medical, and other variables.
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Questionnaires

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25 Questionnaires at all clinical visits include the Short-Form 36 Health Survey (SF-36) and the
26 Hospital Anxiety and Depression Scale (HADS), as well as other questionnaires to assess
27 depressive or disease-related symptoms and sleep qualities. Raw data of these questionnaires
28 have been collected and are available for further analysis.
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Echocardiography

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36 Cardiac chamber geometry and function were assessed via state-of-the-art transthoracic
37 echocardiography. 2D-, Doppler- and M-mode echocardiography were performed following
38 standardized protocols according to current guidelines and are digitally archived.
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Physical functioning and exercise capacity spiroergometry

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45 For the assessment and interaction of vascular disease, cardiovascular risk and biomarkers, an
46 assessment of physical functioning/exercise capacity was essential. Complementary to
47 questionnaires on physical activity, physical fitness and symptom-limited cardiopulmonary
48 exercise capacity was assessed by spiroergometry, six-minute walk test [14], and handgrip
49 measurements [15,16].
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Pulse wave analysis and ECG

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Pulse wave analysis (PWA) and pulse wave velocity (PWV) measurements were performed according to the expert consensus published by Laurent et al [17]. Additionally, a 12-lead ECG was performed in every patient.

Ultrasound of the abdomen

Routine abdominal ultrasound examinations with special attention to liver and kidney characteristics (dimensions, texture, abnormalities) were carried out at each visit.

Carotid ultrasound examinations

Carotid ultrasound examinations were performed in supine position on the right and left side. The intima/media thickness of the left and right common carotid artery is measured [18].

Bone density and body composition measurements

Regular DXA (dual energy X-ray absorptiometry) measurements of bone density at the spine, hip and whole-body density include body composition and trabecular bone score (TBS) [19].

Laboratory and functional metabolic measurements

Sample acquisition includes serum, plasma, saliva and urinary as well as stool samples. Routine tests include liver and kidney function and electrolytes, blood counts, hormonal and metabolic data including lipid profiles and urinary analyses. A standardized oral glucose tolerance tests (oGTT, including insulin and c-peptide) was performed. These materials were collected at each visit for immediate analysis as well as biobanking (including samples of function tests).

Biobanking

Serum, EDTA and citrate plasma, whole blood and cell pellets, spot urine, and saliva were collected at each study visit. They were immediately aliquoted and stored at the biobank of the Medical University of Graz (Austria) at – 80 °C [20]. Biobanking guarantees an accurate description of the sample collection and sample handling according to the STROBE-ME recommendations [13]. Biospecimen-derived measurements adhere to the European guidelines [European guideline issued by the Council of Europe (www.coe.int)].

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Autoantibody phenotyping

Routine measurement of thyroid autoantibodies and autoimmune parameters for celiac disease were performed in all participants. Further exploratory autoantibody detection (endocrine receptors, various epitopes) was done using assays (patent 20170276675) together with the Charité – Universitätsmedizin Berlin, Germany.

Metabolic phenotyping by Nuclear Magnetic Resonance (NMR)

(Un)targeted metabolomic analysis of 1012 baseline serum and urine samples was performed by NMR spectroscopy and state-of-the art chemoinformatics [21–23]. Metabolite and lipoprotein concentrations were determined using Topspin and the Pre-Clinical Screening and In Vitro Diagnostics Research (IVDr) package of Bruker.

Genome-wide characterisation of cohort

A whole genome global screening array (GSA) with nearly 700k specific single nucleotide polymorphisms (SNPs, Illumina, in cooperation with the Human Genotyping Facility at the Erasmus University Rotterdam, NL) is available from all cohort participants.

Optometric phenotyping

In addition, a large part of the cohort has been assessed by non-mydratic retinal photography since 2015. Ophthalmologic examinations comprised undiluted fundus photography and optical coherence tomography including the assessment of retinal vessel diameters.

Data monitoring and quality assurance

All incoming data are checked by the study staff for completeness and plausibility and are entered into an electronic data capture system (OpenClinica; www.openclinica.com), specifically adapted for this project. Additional validation processes such as cross-validation with i.e. external independent validation in samples were regularly performed. External monitoring by certified clinical monitors has been done in 100 % of the baseline study records. Adequate external monitoring of the follow-up data is regularly ongoing.

Patient and Public Involvement

No patient involved.

Findings to date

A descriptive overview of the cohort including the cardiovascular, endocrine, and metabolic risk profile is given in table 1. More female (55 %) than male (45 %) persons were included in the study. The mean age of the participants is 57 ± 8 years and the mean BMI is 26.5 ± 4.5 kg/m². A majority of 59 % of the examined persons has a BMI greater than 25 kg/m². Although asymptomatic, only 26 % of the study population is considered to be at low cardiovascular risk according to the Framingham “Systematic COronary Risk Evaluation” (SCORE) [3], while 38 % show intermediate risk, 26 % show high risk, and 1 % show very high risk. Due to some missing biomarkers at the baseline visit, we were not able to calculate the cardiovascular risk in 9 % of the study population using the Framingham Score. The most common risk factor is hyperlipidaemia (42 % of the population), followed by arterial hypertension (33 % of the population), diabetes mellitus type 2 (T2DM, 18 % of the population), and active smoking (15 % of the population). A rather high number, nearly a third of the study population, has been identified as pre-diabetics based on oGTT data.

The purpose of this manuscript is to describe the study cohort and to give an overview of the baseline characteristics. These are reported in detail including additional information on used materials and methods in the supplementary file (Tables S1 – S10). The central figure 3 shows an R-correlation plot of most measured biomarkers at baseline. In this plot, biomarkers are not grouped (for example in organ systems); instead, clusters of high correlations were formed (blue indicates a positive correlation, red indicates a negative correlation, and white indicates no correlation) [24]. Thereby, associations between different biomarkers of different organ systems can be revealed which may further serve as a basis for a multi-disciplinary in-depth analysis.

Such analyses already identified the so far unknown correlation between IGF1 receptor auto-antibodies with body composition and height as presented at the European Congress of Endocrinology in Barcelona, Spain 2018 [25]. Another preliminary finding revealed a correlation between diabetes status and echocardiographic parameters of the diastolic heart function as presented at the Congress of the European Association for the Study of Diabetes in Barcelona, Spain 2019 [26]. These observations were in line with previous findings of another research group [27]. Furthermore, data from the genome-wide association study of the BioPersMed cohort were already used as control data for a large keratoconus genomic study in cooperation with researchers from the United Kingdom and the Netherlands [28], for

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3 the analysis of allelic determinants with a reported association to 25(OH)D levels and their
4 influence on vitamin D [29], and to identify novel biomarkers for non-alcoholic fatty liver
5 disease [30] in cooperation with local researchers.
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Strengths and limitations

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12 The main strength of the BioPersMed cohort is the joint evaluation of both cardiovascular and
13 metabolic phenotyping including a broad spectrum of diagnostic, imaging, and functional
14 tools. This assures comprehensive biomedical and scientific dimensions of the project within
15 high-end diagnostic and analytical parameters and biomarkers. Second, biobanking with a
16 large number of samples including serum/plasma and blood cells, urine, saliva and stool at
17 each visit that have been aliquoted and stored upon high quality certification rules enabling a
18 prospective view on candidate biomarker in the context of a large longitudinal cohort, where
19 specific approaches can be predefined. In addition, a specifically adapted large electronic
20 database (OpenClinica; www.openclinica.com) assures the quality of data entry and delivery
21 as well as the validity and reliability of biomarkers analyses in the BioPersMed cohort together
22 with a continuous data monitoring.
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33 A potential weakness of this study is the wide time range of a prolonged recruitment due to
34 logistic reasons between 2010 and 2016. This results in a prolonged follow-up period of study
35 participants between 4 and 10 years. Second, after a thorough standardized baseline
36 phenotyping, of the BioPersMed cohort this phenotyping has been expanded by additional
37 diagnostic parameters at either later points in time (e.g. non-mydratic fundoscopy) or in
38 subpopulations. Although some of these biomarkers are not available for the complete
39 duration of the entire cohort, cross-sectional analysis of a considerable number of participants
40 can already be performed with these data sets and will be available for longitudinal
41 comparison of follow-up visits thereafter.
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Collaboration

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53 The design of the BioPersMed study, data management, biobanking and data analyses are
54 compliant to the STROBE, STROBE-ME and STREGA recommendations. Collaboration in data
55 analysis and publications will be welcome through specific research proposals sent to the
56 BioPersMed investigators listed as corresponding authors of this manuscript. If desired,
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retrospective analyses can be performed because all data are recorded as raw data and more than 300,000 samples of blood, serum, and urine are stored in the biobank.

Further Details

Data sharing statement

Data are available upon request. To get access, a proposal must be submitted to the investigators listed as the corresponding authors of this manuscript (*barbara.obermayer@medunigraz.at* and *albrecht.schmidt@medunigraz.at*). Additional information can be obtained via the Research Management of the Medical University of Graz (*tanja.ball@medunigraz.at*)

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Contributorship statement

BP, TP, AS, and BOP designed and supervised the study. AS and BOP administrated the study. CH, EK, NV, BOP, and AS wrote the manuscript. EK, CC, IM, MUM, TG, BH, AL, CR, NV, KA, MW, EPK, NT, GS, and AS investigated participants and collected data. CH, EK calculated the statistics performed principal data analysis. AT, NS, NV, TM, AS, AW, AZ, TK, and RS contributed with data analysis and the allocation of resources. All authors reviewed the manuscript

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Conflict of interest:

The authors declare no conflict of interest.

For peer review only

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Figures

Figure 1

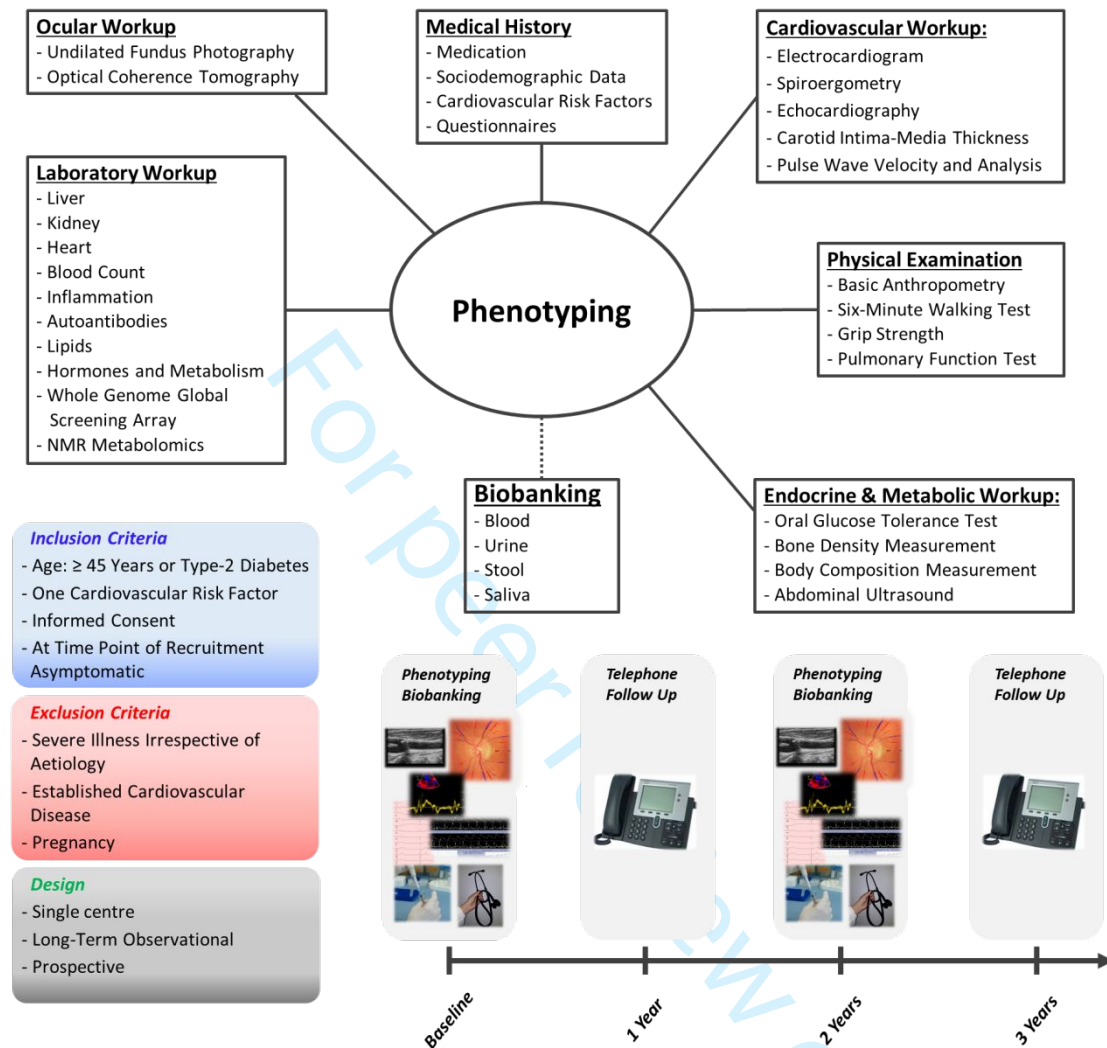


Figure 1

Illustration of the comprehensive phenotyping and biosampling of the BioPersMed Cohort. Follow-up visits are performed according to a tight pre-planned schedule, including reminder-phone calls by study nurses. At baseline, every participant received a patient’s diary for documentation of medical events (source data).

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Figure 2

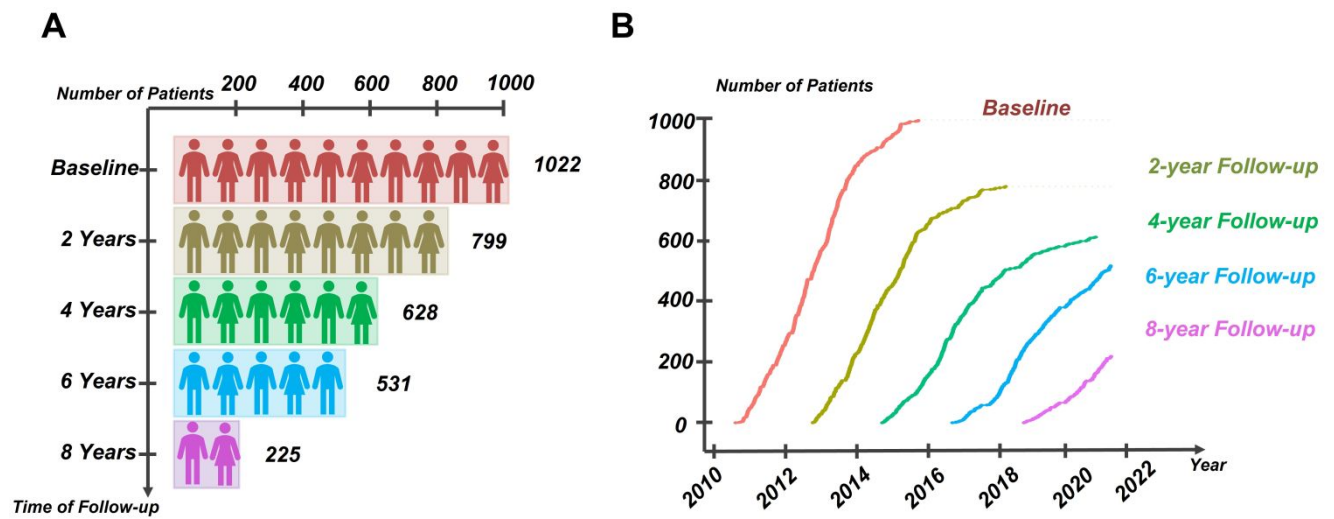


Figure 2

The recruitment status over time and the follow-up of the participants' phenotyping in the BioPersMed Cohort. A: Absolute number of participants who have completed various on-site follow-ups until 1st September 2021. B: Timeline of recruitment and follow-up processes.

Figure 3

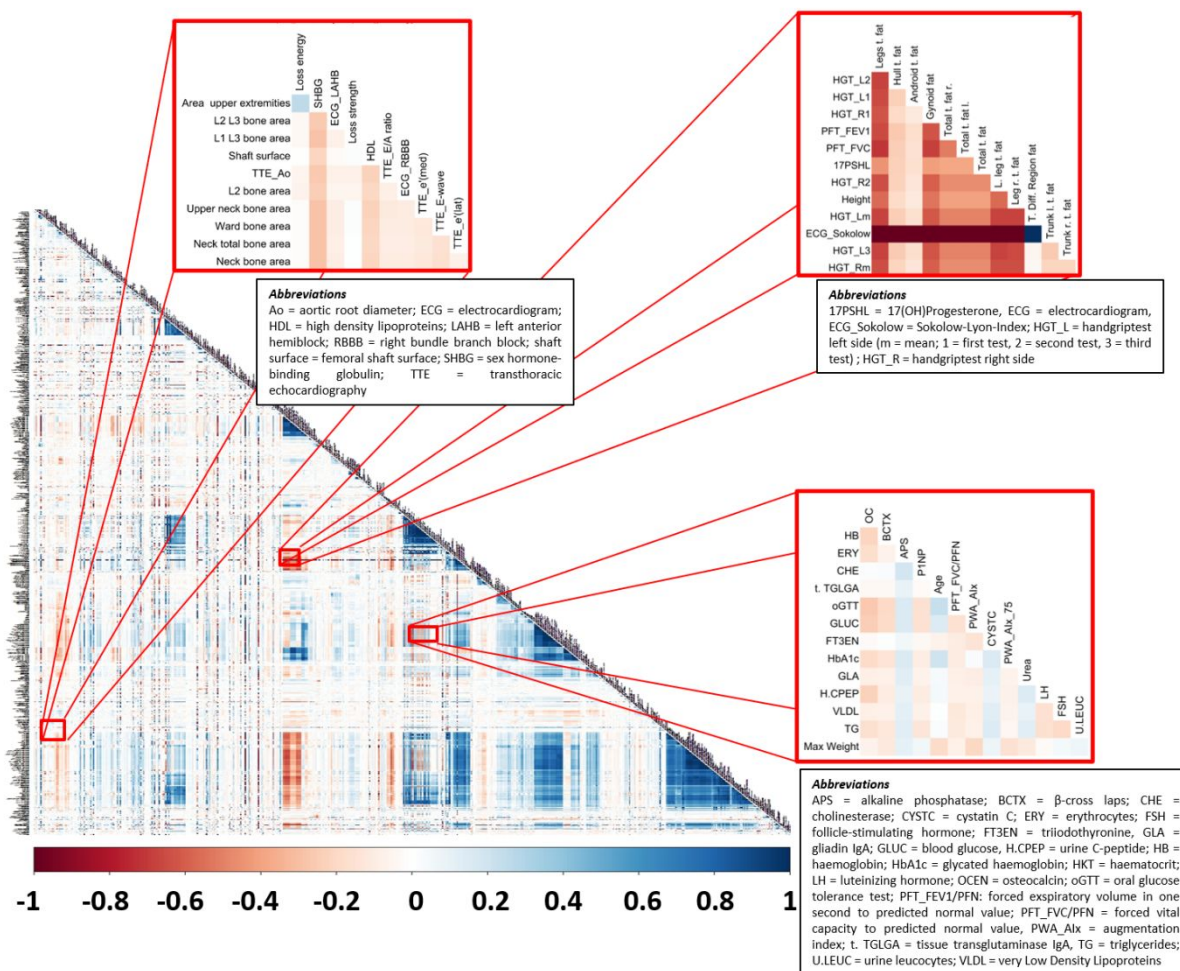


Figure 3

R-plot of the analysed data set. Biomarkers are not grouped; instead, random clusters of correlations are formed. Blue zones indicate positive correlations, red zones indicate negative correlations. As an example, zones of strong correlations are zoomed and biomarkers are depicted.

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Tables**Table 1**

Table 1: Cardiovascular, endocrine, and metabolic risk profile of the BioPersMed cohort (n=1022)		
	Number (N total =1022)	%
Sex		
Men	454	45
Women	568	55
Smoking		
Active smoker	154	15
Former smoker	326	32
Non-smoker	506	49
unknown	36	4
Framingham SCORE*		
Low Risk (< 3 %)	263	26
Intermediate Risk (3 – 4 %)	390	38
High Risk (5 – 9 %)	267	26
Very High Risk (≥ 10 %)	9	1
unknown	93	9
Diabetes status**		
NG	390	38
PreD	315	31
T2DM	181	18
unknown	136	13
Medical history***		
Arterial Hypertension	341	33
Hyperlipidemia	434	42
Stroke / TIA	7	1
Antihypertensive drugs per patient		
No antihypertensive drug	670	65
One antihypertensive drug	154	15
Two antihypertensive drugs	118	12
More than two antihypertensive drugs	80	8
Lipid-lowering drugs per patient		
No lipid-lowering drug	876	85
One lipid-lowering drug	139	14
More than one lipid-lowering drugs	7	1
Antidiabetic drugs per patient		
No antidiabetic drug	844	82
Dietetic treatment	119	12
One antidiabetic drug	54	5

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More than one antidiabetic drugs	5	1
Age		
< 55 years	437	43
55 – 65 years	382	38
66 – 75 years	178	17
> 75 years	25	2
Body mass index		
< 18.5 kg/m ²	6	1
18.5 – 25.0 kg/m ²	408	40
> 25.0 kg/m ²	608	59

Health status - abbreviations

NG = normoglycemia; T2DM = type 2 diabetes mellitus; PreD = prediabetes ; TIA = transient ischemic attack ; * whenever present, a direct categorization due to comorbidities was performed; **Diabetes status based on oGTT results; *** Hyperlipidaemia was assumed based on total cholesterol > 200 mg/dL or the use of a lipid-lowering drug

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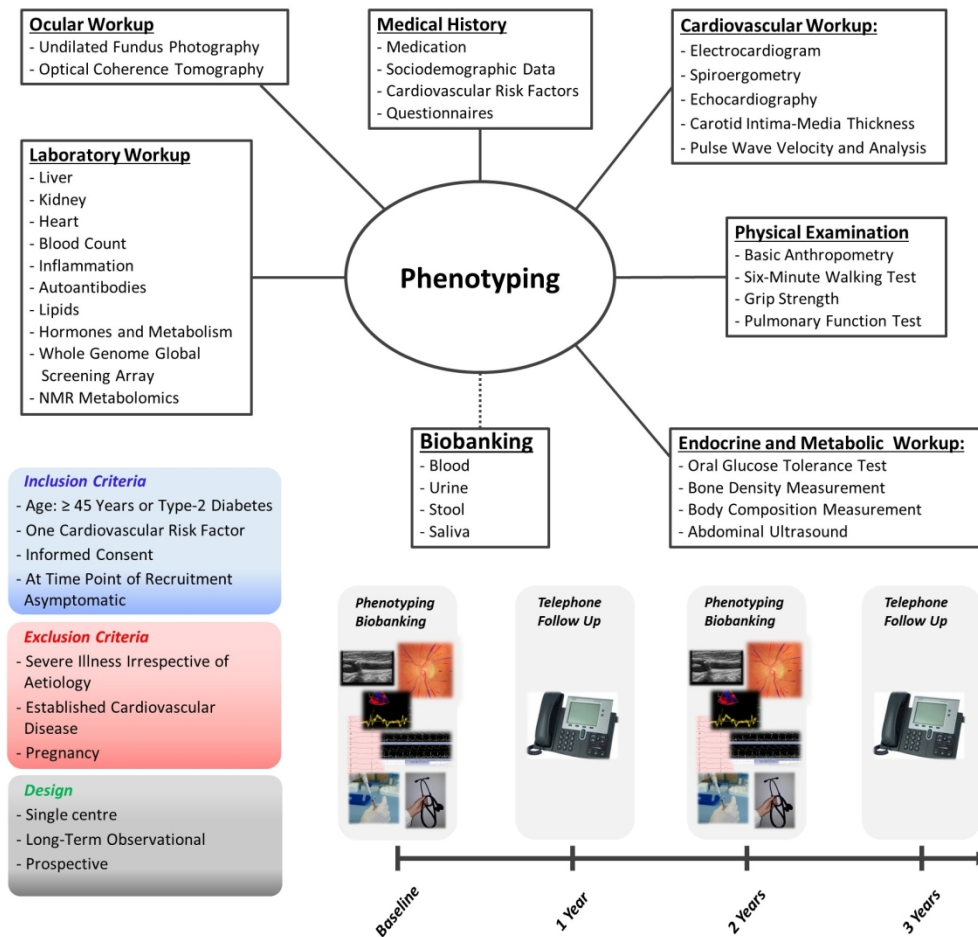


Figure 1

Illustration of the comprehensive phenotyping and biosampling of the BioPersMed Cohort. Follow-up visits are performed according to a tight pre-planned schedule, including reminder-phone calls by study nurses. At baseline, every participant received a patient's diary for documentation of medical events (source data).

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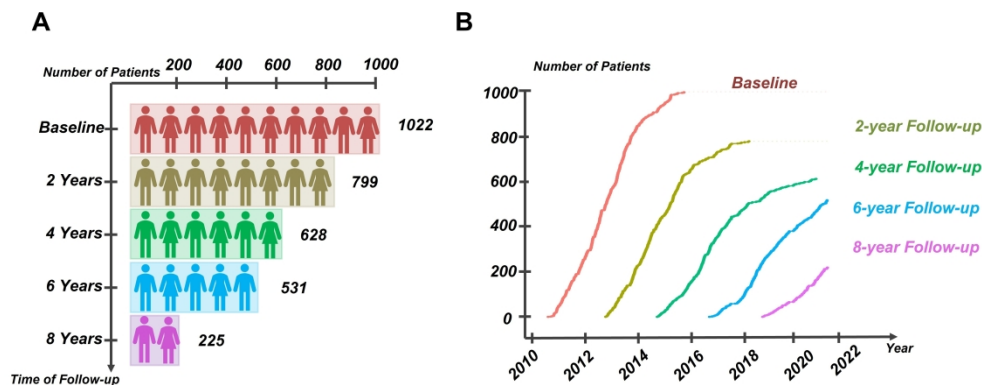
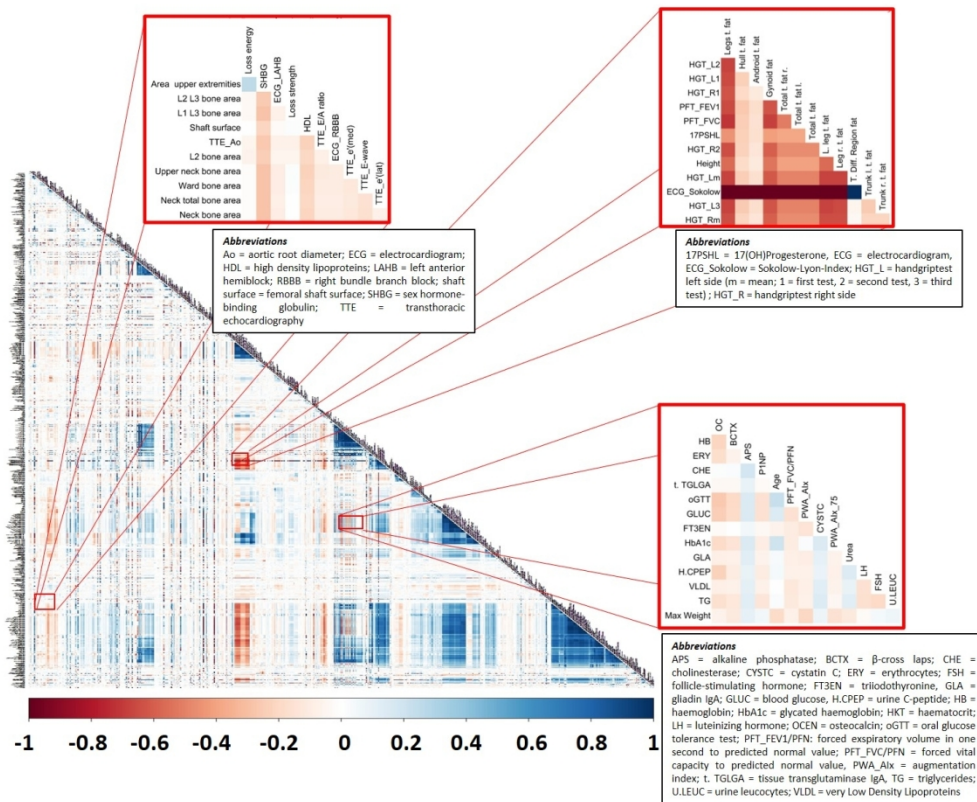


Figure 2

The recruitment status over time and the follow-up of the participants' phenotyping in the BioPersMed Cohort. A: Absolute number of participants who have completed various on-site follow-ups until 1st September 2021. B: Timeline of recruitment and follow-up processes.

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R-plot of the analysed data set. Biomarkers are not grouped; instead, random clusters of correlations are formed. Blue zones indicate positive correlations, red zones indicate negative correlations. As an example, zones of strong correlations are zoomed and biomarkers are depicted.

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Supplements – Methods and baseline characteristics

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1. Echocardiography

For all cardiac ultrasound examinations, a GE vivid E9 device (GE Healthcare, United Kingdom) with a phased array transducer (GE M5s) was used. During the echocardiographic examination, patients were positioned in the left lateral decubitus position. An ECG tracing was recorded during the whole examination. Loops of 3 cardiac cycles of the parasternal long axis view, the parasternal short axis view (on the level of the aortic valve, the mitral valve, the papillary muscles, and the apex), the apical four chamber view, the apical two chamber view, the apical three chamber view, the apical five chamber view, and the subcostal view were recorded. Additionally, colour Doppler loops of all mentioned echocardiographic angulations were recorded for the semi-quantitative assessment of a regurgitation or stenosis. Doppler measurements of the transmitral flow, the transaortic flow, the tricuspid regurgitation flow, and the pulmonic venous flow were recorded as images. Cardiac chambers were quantified by the measurement with the M-Mode technique or as a measurement within the two-dimensional image. M-mode and Doppler images were recorded at a sweep speed of 100 mm/s. The colour Doppler measurements were recorded with a Nyquist limit of 65 ± 5.5 cm/s. For the calculation of the left ventricular mass, the modified Devereux formula [1] was used, for the calculation of the left atrial volume, the biplane formula was used. The body surface area was calculated from height and weight using the DuBois formula. Table S1 gives an overview of all assessed parameters of the baseline-examination.

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Table S1: Echocardiographic parameters of the baseline cohort (N=1022)			
	Method	Mean \pm SD	Unit
Left Ventricle			
IVS <i>end diastolic</i> - PLAX	M-Mode	10.5 \pm 1.9	mm
PW <i>end diastolic</i> - PLAX	M-Mode	10.6 \pm 2.0	mm
LVD <i>end diastolic</i> - PLAX	M-Mode	47.4 \pm 5.9	mm
LVM	Calculated (modified Devereux formula)	184.5 \pm 61.3	g
LVMi	Calculated (LVM indexed to BSA)	96.8 \pm 26.2	g/m ²
LVD <i>end systolic</i> PLAX	M-Mode	29.8 \pm 4.8	mm
LVD apical major axis <i>end diastolic</i> - A4CH	2D	76.1 \pm 8.1	mm
LVD apical minor axis <i>end diastolic</i> - A4CH	2D	43.6 \pm 5.5	mm
LVV <i>end diastolic</i> - A4CH + A2CH	2D (Simpson's biplane method)	91.4 \pm 26.9	mL
LVV <i>end systolic</i> - A4CH + A2CH	2D (Simpson's biplane method)	34.0 \pm 13.1	mL
EF	Calculated (Simpson's law)	63.8 \pm 5.8	%
MAPSE medial annulus - A4CH	M-Mode	0.92 \pm 0.16	cm
MAPSE lateral annulus - A4 CH	M-Mode	1.54 \pm 0.26	cm
Right Ventricle			
TAPSE - A4CH	M-Mode	2.31 \pm 0.09	cm
Left Atrium			
LA minor axis <i>end systolic</i> - PLAX	PLAX	38.4 \pm 5.7	mm
LA major axis <i>end systolic</i> - A4CH	2D	48.0 \pm 6.0	mm
LAA <i>end systolic</i> - A4CH	2D	17.6 \pm 3.6	cm ²
LAA <i>end systolic</i> - A2CH	2D	17.9 \pm 3.6	cm ²
LAV <i>end systolic</i> - A4CH + A2CH	Calculated (biplane method)	56.1 \pm 16.6	mL
LAVI <i>end systolic</i> - A4CH + A2CH	Calculated (LAV indexed to BSA)	29.8 \pm 7.9	mL/m ²
Doppler Measurements			
AV-Vmax	CWD	1.2 \pm 0.3	m/s
Peak E-wave velocity	PWD of transmitral flow	70.0 \pm 16.1	cm/s
Peak A-wave velocity	PWD of transmitral flow	67.2 \pm 18.1	cm/s
Mitral valve deceleration time	PWD of transmitral flow	221 \pm 56	ms
E/A ratio	PWD of transmitral flow	1.10 \pm 0.38	
Duration A-wave	PWD of transmitral flow	132 \pm 37	ms
Duration IVRT	PWD of transmitral flow	115 \pm 20	ms
Duration aortic flow	PWD of transmitral flow	276 \pm 37	ms
Duration E-wave to A-wave	PWD of transmitral flow	447 \pm 48	ms
Peak E-wave velocity (<i>valsalva</i>)	PWD of transmitral flow under <i>valsalva</i>	62.2 \pm 16.0	m/s
Peak A-wave velocity (<i>valsalva</i>)	PWD of transmitral flow under <i>valsalva</i>	63.3 \pm 18.6	m/s
Mitral valve deceleration time (<i>valsalva</i>)	PWD of transmitral flow under <i>valsalva</i>	241 \pm 67	ms
e' velocity (medial annulus)	TDI-PWD	7.6 \pm 2.2	cm/s
e' velocity (lateral annulus)	TDI-PWD	10.4 \pm 3.0	cm/s
E/e' <i>mean</i>	Calculated	8.2 \pm 2.4	
Tricuspid regurgitation systolic jet velocity	CWD	2.25 \pm 0.27	m/s
Pulmonary vein systolic velocity	PWD of pulmonary venous flow	56.7 \pm 12.5	cm/s
Pulmonary vein diastolic velocity	PWD of pulmonary venous flow	42.6 \pm 10.1	cm/s
Pulmonary vein S/D ratio	PWD of pulmonary venous flow	1.37 \pm 0.38	
Flow propagation time	CD M-Mode of transmitral flow	68.6 \pm 31.0	cm/s
Valves			
Semiquantitative evaluation of regurgitation	CD		

Echocardiographic angulations and Doppler measurements - abbreviations:

PLAX = parasternal long axis view; A4CH = apical four chamber view; A2CH = apical two chamber view; CWD = continuous wave Doppler; PWD = pulsed wave Doppler; TDI = tissue Doppler imaging; CD = colour Doppler

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Cardiac structures - abbreviations

IVS = interventricular septum; PW = posterior wall; LVD = left ventricular diameter; LVM = left ventricular mass; LVMI = left ventricular mass index; LVV = left ventricular volume; EF = ejection fraction; MAPSE = mitral annulus plane systolic excursion; TAPSE= tricuspid annulus plane systolic excursion; LA = left atrium; LAA = left atrial area; LAV = left atrial volume; LAVI = left atrial volume index; IVRT = isovolumetric relaxation time

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2. Physical functioning and exercise capacity spiroergometry

The whole study cohort underwent several examinations assessing the cardiac, pulmonary, and exercise capacity. Firstly, a blood pressure measurement with the Boso-medicus uno (Bosch + Sohn GmbH, Germany) was performed in the seating position after 10 minutes of rest. During the diagnostic workup, a spirometry was performed ahead of ergometry. Symptom limited cardiopulmonary exercise testing on a bicycle ergometer starts at a workload of 20W, followed by a stepwise 20W increment every 2 min. Criteria for discontinuation of the exercise test are defined as recommended by the European Society of Cardiology [2]. A standard 12-lead ECG continuously monitors heart rate, ST-segment changes, and arrhythmias. Blood pressure is recorded at rest and then every 2 min. Ventilatory exchange, oxygen uptake (VO_2), and other cardiopulmonary variables are acquired by averaging breath-by-breath measurements over 10 s intervals. Peak heart rate and workload is recorded immediately upon the end of exercise. Peak VO_2 is defined as the maximum value of the last three 10 s averages during exercise and anaerobic threshold is detected using the V-Slope method [3], based on standard operating procedures for spiroergometry. Finally, participants reported the degree of exhaustion expressed as the level of shortness of breath according to a Borg category-ratio scale (0 – 10 points).

After a period of at least 15 minutes rest, a six-minute walk test was performed. Initially, resting blood pressure was assessed; then patients were instructed to walk for six minutes as far as possible without running or jogging. The Borg category-ratio scale (0 – 10 points) was assessed after the six-minute walking test too.

Finally, a handgrip test using a Jamar® Hydraulic Hand Dynamometer (Performance Health UK, United Kingdom) was performed to measure grip strength of both hands. The participant was tested in the seated position with elbow flexed at 90 degrees without touching the trunk.

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The handle position was adjusted to fit the participant's hand. The participant was asked to squeeze the dynamometer at maximal effort for three trials, with a 30-second break between each trial. The average of three trials was calculated for data analysis. Table S2 gives an overview on the single assessed physical functioning and exercise capacity related biomarkers.

Table S2: Baseline parameters of spirometry, spiroergometry, 6-minute walking test and hand-grip strength test			
	Male	Female	Unit
Blood pressure measurement (N=1022)			
SBP – seating position	141 ± 19	133 ± 20	mmHg
DBP – seating position	88 ± 11	85 ± 11	mmHg
Spirometry (N=1022)			
FVC	4.1 ± 0.9	3.0 ± 0.6	L
PNV (FVC)	4.6 ± 0.5	3.1 ± 0.4	L
FVC / PNV	90.3 ± 16.1	97.1 ± 17.9	%
FEV1	3.3 ± 0.8	2.4 ± 0.5	L/s
PNV (FEV1)	3.5 ± 0.4	2.5 ± 0.4	L/s
FEV1 / PNV	93.2 ± 20.0	94.3 ± 19.7	%
FEV1 / FVC	102.3 ± 17.6	102.3 ± 19.8	%
Spiroergometry (N=1022)			
Wmax	158 ± 36	111 ± 23	W
SBP – resting	125 ± 19	118 ± 19	mmHg
DBP – resting	83 ± 12	78 ± 11	mmHg
SBP - maximal	206 ± 23	189 ± 25	mmHg
DBP - maximal	90 ± 16	89 ± 18	mmHg
HR – resting	77 ± 13	79 ± 13	/min
HR - maximal	143 ± 20	147 ± 19	/min
Peak VO ₂	23.9 ± 7.0	20.5 ± 6.4	mL/kg/min
Anaerobic threshold	107 ± 41	79 ± 25	W
Borg-CR10	3.9 ± 2.0	3.9 ± 2.0	
Six-minute walk test (N=980)			
SBP – resting	139 ± 18	133 ± 20	mmHg
DBP – resting	88 ± 11	85 ± 11	mmHg
SBP - maximal	147 ± 22	141 ± 22	mmHg
DBP - maximal	89 ± 11	88 ± 12	mmHg
HR – resting	73 ± 14	73 ± 11	/min
HR - maximal	80 ± 16	82 ± 14	/min
Distance walked	521 ± 90	501 ± 91	m
Prematurely terminated	3 / 446	4 / 559	N / total
Borg-CR10	0.4 ± 0.8	0.6 ± 1.2	
Hand-grip test (N=680)			
Right arm	96 ± 20	59 ± 13	lbs
Left arm	90 ± 20	54 ± 13	lbs

spiroergometry, six-minute walk test & hand-grip test - abbreviations

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1
2
3 FVC = forced vital capacity; PNV = predicted normal value; FEV1 = forced expiratory volume (1 second); Wmax = maximal
4 workload; SBP = systolic blood pressure; DBP = diastolic blood pressure; HR = heart rate; VO₂ = oxygen uptake; Borg-CR10 =
5 Borg category-ratio scale
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3. Pulse wave analysis and ECG

Measurements of pulse wave analysis (PWA) and pulse wave velocities (PWV) were performed with the SphygmoCor® device (Atcor Medical, Australia). The measurement takes place in supine position after a minimum of 10 to 15 min rest in a quiet, temperature-controlled room after measuring the blood pressure on the right and left side [4]. Central BP measurements were recorded noninvasively by applanation tonometry.

PWV (carotid–femoral), which reflects arterial stiffness, were measured by using the foot-to-foot velocity method. The waveforms were recorded transcutaneously at the right common carotid artery and the right femoral artery. In addition, the augmentation index was determined by applanation tonometry on the central pressure waves measured in the right radial artery according to previous recommendations [5]. The SphygmoCor device quality index (QI), which represents waveform reproducibility, was checked after each pulse wave analysis (PWA) and the measurements are repeated in case of an $QI < 80$. As an instant quality check for the measurement of the PWV, the time difference between the ECG-signal and the signal from the recording sites was suggested to have a $SD \leq 10\%$ of the mean value. All measurements of PWA and PWV were performed in duplicate.

Briefly after the pulse wave analysis, a 12-lead resting ECG was performed. An overview of the ECG and PWA related biomarkers is shown in table S3.

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Table S3: Baseline parameters of 12-lead ECG and pulse wave velocity/analysis		
	Mean \pm SD	Unit
12-lead ECG (N=1019)		
HR	64.4 \pm 11.1	/min
PR-Interval	159.3 \pm 24.9	ms
QRS-duration	95.8 \pm 15.9	ms
QTc-time	418.6 \pm 27.6	ms
PWA (N=993)		
SBP	126 \pm 18	mmHg
DBP	86 \pm 11	mmHg
AP	12.2 \pm 7.6	mmHg
Alx	28.2 \pm 11.4	%
Alx(HR75)	23.9 \pm 11.3	%
PP	39.9 \pm 13.1	mmHg
PWV (N=963)		
Carotid-femoral PWV	7.4 \pm 2.1	m/s

12-lead-ECG and pulse wave analysis – abbreviations

HR = heart rate; SBP = systolic blood pressure; DBP = diastolic blood pressure; AP = augmentation pressure; Alx = augmentation index; Alx(HR75) = augmentation index normalized to the heart rate of 75 beats per minute; PP = pulse pressure

4. Ultrasound of abdomen

An ultrasound of the abdomen was performed using the GE vivid E9 ultrasound device with a curved array transducer (GE CI5). The dimensions of the spleen were assessed, and the echogenicity of the liver was evaluated in order to identify signs of liver-steatosis. The findings of the abdominal ultrasound are summarized in table S4.

	Number (N total = 982)	%
Ultrasound of abdomen		
Normal	626	64
Blunted liver edge	32	3
Irregular liver surface	2	< 1
Ascites	1	< 1
Signs of steatosis	321	33
Dimension of spleen		
	Mean ± SD	Unit
Longitudinal	101.8 ± 14.4	mm
Transversal	36.6 ± 7.7	mm

5. Carotid ultrasound examinations

Carotid ultrasound examination was performed with a GE vivid E9 ultrasound device with a linear array transducer (GE 9L). Each measurement includes both the near-wall and far-wall data respectively for the right and left carotid arteries [6]. The common carotid intima-media thickness (CIMT) was calculated by an automated measurement program by GE at more than 200 points distal from the bulbous. Sonographers are taking images showing the maximum thickness of a particular site. Detected plaques at specific sites are included in the maximum CIMT measurement. Table S5 gives an overview on the biomarkers of the carotid ultrasound examination.

Table S5: Baseline parameters of carotid ultrasound		
	Mean \pm SD	Unit
CIMT		
Right side	0.73 \pm 0.16	mm
Left side	0.73 \pm 0.17	mm
Presence of calcified plaques or stenosis	Number (N=1021)	%
Right side	117	11
Left side	106	10

Carotid ultrasound – abbreviation:

CIMT = carotid intima-media thickness

6. Bone density and body composition

Bone imaging via 2D bone density measurements and body composition measurements were performed using Lunar iDXA (General Electrics, USA) at each visit including measurements at the lumbar spine, the hip and whole-body sites. Thorough body composition, automated comparison to prior measurements and an additional trabecular bone score (TBS, as a measure of bone texture correlated with bone microarchitecture, software by iDEXA solutions, Switzerland [7]) has been documented. Table S6 shows an overview on the parameters of the bone density measurements.

	Male (N=341)	Female (N=448)	%
Bone mineral density			
normal BMD	325	412	93
Osteopenia or osteoporosis	16	36	7
Body Composition	Mean ± SD	Mean ± SD	Unit
Total BMD	1.28 ± 0.11	1.13 ± 0.12	g/cm ²
Total BMC	3132 ± 392	2293 ± 336	g
Total T-Score	0.78 ± 1.09	0.48 ± 1.14	STD
Total Z-Score	0.63 ± 1.04	1.01 ± 0.94	STD
Total Tissue mass	84397 ± 13301	68502 ± 13681	g
Total Bone mass	3194 ± 243	2180 ± 181	g
L1-L4 BMD	1.25 ± 0.19	1.14 ± 0.17	g/cm ²
L1-L4 BMC	82.2 ± 16.4	62.4 ± 12.5	g
L1-L4 T-Score	0.09 ± 1.52	-0.43 ± 1.36	STD
L1-L4 Z-Score	0.14 ± 1.51	0.29 ± 1.19	STD

Bone density measurements – abbreviations:

BMD = Bone mineral density in [g/cm²], BMC = Bone mineral content [g], L1-L4 = region vertebrae L1-L4, T-Score= standard deviations from mean BMD level of young adults, Z- or T-Score STD = standard deviations from age adjusted mean BMD level

7. Laboratory assessment and measurements

Routine clinical laboratory parameters such as blood count and coagulation, serum electrolytes, kidney and liver function, HbA1c, fasting c-peptide, insulin as well as fasting blood glucose followed by a standardized oral glucose tolerance test (oGTT, at 30, 60 and 120 minutes after 75mg glucose load) with consecutive insulin and c-peptide measurements, high-sensitive c-reactive protein (hsCRP), lipids (total cholesterol, high-density and low-density lipoproteins (HDL and LDL), lipoprotein a (LP(a), triglycerides (TG), high-sensitive troponin T (hsTropT) and cardiac parameters such as creatine kinase (CK, CK-MB), as well as urinary proteins, electrolytes, cell count and c-peptide are measured in all participants at each study visit. The baseline measurements are presented in table S7.

In addition to the oGTT and glucose metabolism monitoring, a large panel of hormonal and metabolic analyses are performed in women as well as in men. These biomarkers include estrogen, testosterone and free testosterone, androstenedione, dehydroepiandrosterone (DHEAS), 17(OH)progesterone, and anti-Müllerian hormone (AMH), saliva hormones (e.g. cortisol), bone metabolism including formation and resorption parameters (osteocalcin, N-terminal propeptide of type 1 procollagen (P1NP), serum-crosslaps (CTX)), calciotropic hormones such as 25(OH)vitamin D and parathyroid hormone (PTH), pituitary hormones (LH, FSH, ACTH) and others.

Routine biochemical parameters (liver, kidney, lipid, electrolyte and other profiles) were determined using a Cobas® Analyzer (Roche Diagnostics, Germany), complete blood counts by Beckmann-Coulter, Germany. Hormonal measurements such as insulin and c-peptide were measured by chemiluminescence on an Advia Centaur system (Siemens Healthcare Diagnostics, Germany). Automated analysers were used to measure hormones: anti-Müllerian hormone (AMH) (Beckmann-Coulter, Germany); testosterone, cortisol, thyrotropin,

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triodothyronine and thyroxine (Siemens Advia Centaur, Germany); sexual hormone-binding globulin (SHBG) (Roche Diagnostics, Germany); estrogen, luteotropic hormone (LH) and follicle-stimulating hormone (FSH) (Triturus, Biomedical Diagnostics, Antwerp, Belgium); 25(OH)vitamin D and bone metabolism parameters such as bone-specific alkaline phosphatase (bALP), osteocalcin (OC), C-terminal telopeptide (CTX) and procollagen type 1 N-terminal propeptide (P1NP) by iSYS, (IDS, UK, or Roche Diagnostics, Germany, respectively). Endocrine and bone biomarkers are shown in table S8, all of them collected in the morning after an overnight fast.

Table S7: Baseline parameters of general, metabolic and cardiovascular laboratory assessment			
	Male (N=444)	Female (N=556)	Unit
Liver	Mean ± SD	Mean ± SD	
ALP	93 ± 32	109 ± 11	U/L
ALT	24.7 ± 31.3	16 ± 7.3	U/L
AST	20.9 ± 17.1	18.1 ± 7.4	U/L
CHE	11059 ± 928	11161 ± 1153	U/L
GGT	44.8 ± 54.2	18 ± 20.1	U/L
LDH	161 ± 19	166 ± 19	U/L
Kidney			
Creatinine	0.98 ± 0.14	0.8 ± 0.13	mg/dL
Creatinine/urine	142 ± 35	130 ± 45	mmol/L
Urea	18.8 ± 1.5	18.1 ± 2.2	mg/dL
Uric Acid	6.3 ± 1.2	4.8 ± 1.2	mg/dl
Proteins			
Albumin	4.6 ± 0.3	4.6 ± 0.3	g/dL
CRP	2.3 ± 4.1	2.5 ± 3.7	pg/mL
Ferritin	158 ± 173	1235 ± 47	ng/mL
Total Protein	7.4 ± 0.4	7.4 ± 0.4	g/dL
Transferrin	2.5 ± 0.3	2.5 ± 0.3	g/L
Electrolytes & Iron			
Calcium/urine	2.41 ± 1.88	2.21 ± 1.69	mmol/L
Chloride	103 ± 2	104 ± 2	mmol/L
Chloride/urine	1489 ± 33	135 ± 41	mmol/L
Iron	132 ± 24	126 ± 22	µg/dL
Magnesium	0.85 ± 0.06	0.85 ± 0.06	mmol/L
Phosphate	2.61 ± 0.75	2.96 ± 0.49	mg/dL
Phosphate/urine	18.6 ± 14.5	15.1 ± 10.8	mg/dl
Potassium	4.1 ± 0.5	4.0 ± 0.3	mmol/L
Total Calcium	2.39 ± 0.09	2.39 ± 0.1	mmol/L
Haematopoiesis			
Haemoglobin	15.1 ± 1.0	13.6 ± 0.9	g/dL
Platelets	174 ± 20	180 ± 18	10 ⁹ /L
TLC	5.77 ± 1.44	5.76 ± 1.76	10 ⁹ /L
Glucose metabolism			
Glucose fasting	99 ± 20	91 ± 17	mg/dL

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Glucose 30 min	158 ± 38	142 ± 36	mg/dL
Glucose 60 min	158 ± 62	128 ± 52	mg/dL
Glucose 120 min	115 ± 57	100 ± 46	mg/dL
Insulin fasting	13 ± 17	10 ± 8	μU/mL
Insulin 30 min	64 ± 54	56 ± 48	μU/mL
Insulin 60 min	94 ± 87	73 ± 68	μU/mL
Insulin 120 min	62 ± 72	51 ± 62	μU/mL
HbA1c	38.6 ± 7.1	37.7 ± 5.8	mmol/mol
Heart			
CK	151 ± 132	130 ± 25	IU
NT-Pro-BNP	79 ± 18	122.66 ± 105.7	pg/mL
PT	105.7 ± 5.0	106.8 ± 5.2	%
Lipid parameters			
Cholesterol	174.8 ± 18.9	181.6 ± 16.6	mg/dL
LDL	130.9 ± 36.5	132.1 ± 33.6	mg/dL
VLDL	19.6 ± 21.9	15.7 ± 2.8	mg/dL
HDL	48.5 ± 16.3	65.1 ± 20.2	mg/dL
LPA	1.37 ± 3.47	1.81 ± 4.16	mg/dL
Triglycerides	144 ± 80	133 ± 26	mg/dL

Laboratory measurements – abbreviations:

ALP = Alkaline Phosphatase; ALT = Alanine Aminotransferase; AST = Aspartate Aminotransferase; CHE = Cholinesterase; CRP = C-Reactive Protein; GGT = Gamma Glutamyl Transferase; LDH = Lactate Dehydrogenase; HB = Haemoglobin; PLT = Platelets; TLC = Total leucocyte count; HbA1c = glycated haemoglobin; CK = Creatine Kinase; NT-Pro-BNP = N-terminal pro brain natriuretic peptide; PT = Prothrombin Time; LDL = Low Density Lipoprotein; VLDL = Very low Density Lipoprotein; HDL = High Density Lipoprotein; LPA = Lipoprotein A

	Male (N=446)	Female (N=557)	Unit
Endocrine parameters	Mean ± SD	Mean ± SD	
17(OH)Progesterone	0.9 ± 0.4	0.4 ± 0.3	ng/ml
ACTH	21.3 ± 11.4	14.7 ± 8.1	pg/ml
Aldosterone	8.7 ± 7.6	9.0 ± 8.1	ng/dl
Androstenedione	2.6 ± 1	1.9 ± 0.9	ng/ml
Cortisol	123.2 ± 36.5	111.9 ± 38.13	ng/ml
Cortisol/saliva	7 ± 4.4	6.4 ± 4.2	ng/ml
DHEAS	1.1 ± 0.7	0.7 ± 0.5	μg/ml
Estradiol	58.4 ± 280.8	49.4 ± 63.8	pg/ml
Free Thyroxine	15.1 ± 2.2	15.1 ± 2.7	pmol/l
Free Triiodothyronine	5 ± 0.5	4.6 ± 0.6	pmol/l
FSH	9.34 ± 8.7	67.2 ± 30.9	mU/ml
Insulin	13.4 ± 16.9	9.9 ± 7.7	mU/l
LH	6.0 ± 5.7	14.2 ± 6.4	mU/ml
Progesterone	0.51 ± 1.18	0.9 ± 4.06	ng/ml
Renin	58.3 ± 106.5	30.9 ± 74.4	μU/ml
SHBG	41.1 ± 42.1	119.5 ± 51.8	nmol / L
Total Testosterone	11.0 ± 3.9	1.4 ± 0.7	pg/ml
TSH	2.0 ± 1.1	2.2 ± 6.3	μU/ml
Bone parameters			
25(OH)Vitamin D	81.0 ± 28.3	83.7 ± 29.8	ng/ml
Crosslaps/serum	0.3 ± 0.12	0.37 ± 0.17	ng/ml

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Osteocalcin	18.8 ± 5.6	23.2 ± 8.7	ng/ml
P1NP	40.4 ± 13.6	49.8 ± 20.4	ng/ml

Laboratory measurement: endocrinology & bone metabolism - abbreviations:

ACTH = Adrenocorticotrophic hormone DHEAS = dehydroepiandrosterone sulfate, FSH = Follicle-stimulating hormone, LH = Luteinizing hormone, SHBG = Sex hormone-binding globulin, TSH = Thyroid-stimulating hormone, thyrotropin, P1NP = procollagen type 1 N-terminal propeptide

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8. Autoantibody phenotyping

Data are available for routine thyroid and gluten autoantibodies, further autoimmune parameters for glucose and bone metabolism, body growth, and cardiovascular risk were analysed using a non-automated bridge assay protocol based on the method from EU patent 20170276675 together with Charité – Universitätsmedizin Berlin, Germany. Gliadin and tissue transglutaminase autoantibodies were measured by an automated Chemiluminescence Immunoassay (CLIA) from IDS (Immunodiagnostic Systems, Boldon, UK), see table S9. More autoantibodies are currently measured for endocrine and potentially cardiovascular receptors and targets of interest.

	Male (N=377)	Female (N=484)	Unit
Laboratory assessment	Mean ± SD	Mean ± SD	
Gliadin IgA	2.2 ± 4.9	0.7 ± 2.1	ng/ml
Gliadin IgG	2.0 ± 11.2	1.6 ± 11.4	ng/ml
Tissue transglutaminase IgA	2.5 ± 0.8	2.2 ± 0.5	ng/ml
Tissue transglutaminase IgG	0.3 ± 0.9	0.3 ± 1.1	ng/ml

Laboratory measurement: autoantibody phenotyping - abbreviations:

IgA = Immunoglobulin A, IgG = Immunoglobulin G

9. Metabolic phenotyping by Nuclear Magnetic Resonance (NMR)

Untargeted metabolomics was done in 1012 serum and urine samples using NMR-based metabolic phenotyping by NMR spectroscopy. Technical measurements were performed on a 600 MHz Avance Neo NMR spectrometer and using 1D CPMG (Carr–Purcell–Meiboom–Gill), NOESY and 2D J-resolved pulse sequences. Data processing: Principle Component Analysis (PCA) and Orthogonal Partial Least Squares Discriminant Analysis (O-PLS-DA) were performed. Metabolite reference chemical shifts from databases and metabolites were cross-checked using reference compounds and Chenomx software package if necessary. Metabolite concentrations were determined using internal/external standards and using the Eretic 2 approach implemented in Topspin [8,9].

10. Genome-wide characterization of cohort

More than 650.000 single nucleotide polymorphisms (SNPs) have been measured in a whole genome global screening array (GSA) in all cohort patients in cooperation with the Human Genotyping Facility (Genetic Lab at the Erasmus University Rotterdam, Netherlands) based on Illumina technologies in 2017. These genetic data, as well as specific genotyping e.g. for primary hypolactasia, are available for genome-wide association analyses and the inclusion in subsequent meta- and mega-analyses. As the results are stored in database, bioinformatical approaches such as diverse R-scripts allow for a fast gene-, pathway- or function-specific selection and export of SNP data. Epigenetic phenotyping is ongoing. The potential of various circulating non-coding RNAs for the diagnosis and prediction of (subclinical) diseases is currently evaluated for diabetes, as well as osteological, metabolic and CVD conditions.

11. Optometric phenotyping

For the ophthalmologic examinations, a Canon CR-2 AF (Canon Europa, Amstelveen, Netherlands) and for the optical coherence tomography (OCT) an OCT Spectralis (Heidelberg Engineering, Heidelberg, Germany) was used. Retinal vessel diameters were assessed with Integrative Vessel Analysis (IVAN software, N. Ferrier, University of Wisconsin, USA) and with the OCT device's inbuilt calliper tool from 12° peripapillary circle scans. Additional OCT scans included a macular volume scan and a high-resolution single scan through the fovea. Measured parameters are described in Table S10.

Table S10: Baseline parameters of optometric phenotyping		
	Mean \pm SD	Unit
Right eye	Number (N=349)	
CRAE6	147.5 \pm 12.5	μm
CRVE6	212.2 \pm 17.7	μm
AVR	0.70 \pm 0.06	
Left eye	Number (N=344)	
CRAE6	148.3 \pm 13.6	μm
CRVE6	212.1 \pm 18.1	μm
AVR	0.70 \pm 0.06	

Optometric phenotyping – abbreviations:

CRAE6: mean diameter of the six biggest arterioles; CRVE6: mean diameter of the six biggest venules; AVR: arterioles-venules ratio

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12. Biobank Storage

Until 2020, samples were transferred immediately after sampling to an adjacent laboratory run by technicians from the local biobank. Sodium citrate samples were centrifuged at 2880g for 15min and the rest of the samples at 3360g for 10min at 4°C. The centrifuged biospecimen were then aliquoted manually. Since November 2013 samples were automatically aliquoted using a pipetting robot (Hamilton Microlab STARlet). After aliquoting, samples were temporarily stored at -20°C. The transport to the final -80°C Biobank Graz storage unit was performed under ambient temperature. Since March 2019 the transport was performed under dry-ice conditions. Since 2020 these processes took place in another laboratory and technicians from the biobank picked up the samples at latest within one hour after sampling. Additionally, timepoints of sampling and execution of single work steps were digitally documented.

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BMJ Open

Cohort profile: "Biomarkers of Personalized Medicine" (BioPersMed) – A single-centre prospective observational cohort study in Graz/Austria to evaluate novel biomarkers in cardiovascular and metabolic diseases

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Complete List of Authors:	<p>Haudum, Christoph; Center for Biomarker Research in Medicine , ; Medical University of Graz, Department of Internal Medicine</p> <p>Kolesnik, Ewald; Medizinische Universität Graz, Department of Internal Medicine and University Heart Center Graz</p> <p>Colantonio, Caterina ; Medizinische Universität Graz, Department of Internal Medicine and University Heart Center Graz</p> <p>Mursic, Ines; Medical University of Graz, Department of Internal Medicine</p> <p>Url-Michitsch, Marion; Medical University of Graz, Department of Internal Medicine and University Heart Center Graz</p> <p>Tomaschitz, Andreas; Medical University of Graz, Department of Internal Medicine and University Heart Center Graz</p> <p>Glantschnig, Theresa ; Medical University of Graz, Department of Internal Medicine and University Heart Center Graz</p> <p>Hutz, Barbara; Medical University of Graz, Department of Internal Medicine</p> <p>Lind, Alice ; Medical University of Graz, Department of Internal Medicine</p> <p>Schweighofer, Natascha; Medical University of Graz, Department of Internal Medicine</p> <p>Reiter, Clemens ; Medical University of Graz, Department of Internal Medicine and University Heart Center Graz; Medizinische Universität Graz, Department of Radiology</p> <p>Ablasser, Klemens; Medical University of Graz, Department of Internal Medicine and University Heart Center Graz</p> <p>Wallner, Markus; Medical University of Graz, Department of Internal Medicine and University Heart Center Graz; Lewis Katz School of Medicine at Temple University, Cardiovascular Research Center</p> <p>Tripolt, Norbert; Medizinische Universität Graz, Department of Internal Medicine</p> <p>Pieske-Kraigher, Elisabeth; Charité University Medicine, Department of Internal Medicine and Cardiology</p> <p>Madl, Tobias; Medical University of Graz, Gottfried Schatz Research Center; BioTechMed</p> <p>Springer, Alexander; Medical University of Graz, Gottfried Schatz Research Center; BioTechMed</p> <p>Seidel, Gerald; Medical University of Graz, Department of Ophthalmology</p> <p>Wedrich, Andreas; Medizinische Universität Graz, Department of Ophthalmology</p>

	Zirlik, Andreas; Medical University of Graz, Department of Internal Medicine and University Heart Center Graz Krahn, Thomas; Maastricht University, Department of Pharmacology and Personalised Medicine Stauber, Rudolf; Medizinische Universität Graz, Department of Internal Medicine Pieske, Burkert; Charité University Medicine, Department of Internal Medicine and Cardiology, Pieber, Thomas; Medical University of Graz, Department of Internal Medicine; Center for Biomarker Research in Medicine Verheyen, Nicolas; Medizinische Universität Graz, Department of Internal Medicine and University Heart Center Graz Obermayer-Pietsch, Barbara; Medical University of Graz, Department of Internal Medicine Schmidt, Albrecht; Medizinische Universität Graz, Department of Internal Medicine and University Heart Center Graz
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BioPersMed Concept Paper

Cohort profile: “Biomarkers of Personalized Medicine” (BioPersMed) – A single-centre prospective observational cohort study in Graz/Austria to evaluate novel biomarkers in cardiovascular and metabolic diseases

Authors:

Haudum CW^{1,2°}, Kolesnik E^{3°}, Colantonio C³, Mursic I¹, Url-Michitsch M³, Tomaschitz A³, Glantschnig T³, Hutz B¹, Lind A¹, Schweighofer N¹, Reiter C^{3,4}, Ablasser K³, Wallner M^{3,10}, Tripolt N¹, Pieske-Kraigher E⁵, Madl T^{6,11}, Springer A^{6,11}, Seidel G⁷, Wedrich A⁷, Zirlik A³, Krahn T^{8,12}, Stauber R⁹, Pieske B⁵, Pieber TR^{1,2}, Verheyen N³, Obermayer-Pietsch B^{1*§}, Schmidt A^{3*§}

[°] and ^{*} - these authors contributed equally to the manuscript

[§] Authors for correspondence and reprint requests

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Affiliations:

¹ Department of Internal Medicine, Division of Endocrinology and Diabetology, Medical University of Graz, Graz, Austria

² Center for Biomarker Research in Medicine (CBmed), Graz, Austria.

³ Department of Internal Medicine and University Heart Center Graz, Division of Cardiology, Medical University of Graz, Graz, Austria

⁴ Department of Radiology, Medical University of Graz, Graz, Austria

⁵ Department of Internal Medicine and Cardiology, Charité University Medicine, Campus Virchow Klinikum and German Heart Center, Berlin, Germany

⁶ Gottfried Schatz Research Center, Chair of Molecular Biology and Biochemistry, Medical University of Graz, Graz, Austria

⁷ Department of Ophthalmology, Medical University of Graz, Graz, Austria.

⁸ Bayer AG, Berlin, Germany

⁹ Department of Internal Medicine, Division of Gastroenterology and Hepatology, Medical University of Graz, Graz, Austria

¹⁰ Lewis Katz School of Medicine, Temple University, Cardiovascular Research Center, Philadelphia, PA, USA

¹¹ BioTechMed-Graz, Graz, Austria.

¹² Department of Pharmacology and Personalised Medicine, Faculty of Health, Medicine and Life Sciences, Maastricht University, Maastricht, Netherlands

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39 **Abstract [244/max 250w]**

40 **Purpose**

41 Accumulating evidence points towards a close relationship between cardiovascular, endocrine
42 and metabolic diseases. The BioPersMed Study (Biomarkers of Personalized Medicine) is a
43 single-centre prospective observational cohort study with repetitive examination of
44 participants in 2-year intervals. The aim is to evaluate the predictive impact of various
45 traditional and novel biomarkers of cardiovascular, endocrine and metabolic pathways in
46 asymptomatic individuals at risk for cardiovascular and/or metabolic disease.

48 **Participants:**

49 Between 2010 and 2016, we recruited 1022 regional individuals into the study. Subjects aged
50 45 years or older presenting with at least one traditional cardiovascular risk factor or manifest
51 type 2 diabetes mellitus were enrolled. The mean age of the participants was 57 ± 8 years, 55
52 % were female, 18 % had T2DM, 33 % suffered from arterial hypertension, 15 % were smokers,
53 42 % had hyperlipidaemia, and only 26 % were at low cardiovascular risk according to the
54 Framingham "Systematic COronary Risk Evaluation" (SCORE).

56 **Findings to date:**

57 Study procedures during screening and follow-up visits included a physical examination and
58 comprehensive cardiovascular, endocrine, metabolic, ocular, and laboratory workup with
59 biobanking of blood and urine samples. The variety of assessed biomarkers allows a full
60 phenotyping of individuals at cardiovascular and metabolic risk. Preliminary data from the
61 cohort and relevant biomarker analyses were already used as control population for genomic
62 studies in local and international research cooperation.

64 **Future plans:**

65 Participants will undergo comprehensive cardiovascular, endocrine, and metabolic
66 examinations for the next decades and clinical outcomes will be adjudicated prospectively.

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Strength and Limitations of this study

- The main strength of the BioPersMed cohort is the joint evaluation of cardiovascular, endocrine and metabolic phenotyping including a broad spectrum of highly innovative diagnostic, imaging and functional tools.
- Biobanking with a large number of samples aliquoted and stored at each visit enables a prospective view on candidate biomarkers in the context of a large longitudinal cohort, where specific approaches can be predefined.
- A specifically adapted large electronic data capture system (OpenClinica; www.openclinica.com) and iterated monitoring assures the quality of data entry and delivery as well as the validity and reliability of biomarkers analyses.
- A potential weakness of this study is the wide time range of recruitment due to logistic reasons between 2010 and 2016, with a prolonged follow-up period of study participants to date between 4 and 10 years.
- Some biomarkers are not available for the complete duration of the entire cohort

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87 Introduction

88 Cardiovascular (CVD) and metabolic diseases (MD) are globally representing the most
89 important cause of disability and premature death [1]. Next to our genetic programming,
90 modern lifestyle, including the use of tobacco, unhealthy nutritional habits, physical inactivity,
91 and psychosocial stress are major risk factors of CVD and metabolic diseases within different
92 age groups not only promoting excess cardiovascular (CV) and metabolic morbidity but
93 ultimately triggering excess mortality [2,3]. In turn, primary prevention of these diseases has
94 the potential to avoid many of related deaths [4]. However, the initial euphoria about a
95 decline of CVD prevalence at the beginning of this century [5] gradually gives way to a sense
96 of frustration. This comes in the light of increasing numbers of type 2 diabetes mellitus (T2DM)
97 [6], a very high lifetime-risk for the development of heart failure with stable incidence over
98 the last decades [7], and a persistent high stroke mortality [8]. In addition, the relevant
99 underdiagnosing and/or undertreatment of patients at high risk for CVD related risk factors
100 remains of utmost important [9]. Therefore, early detection of asymptomatic CV and/or
101 metabolic risk remains a crucial challenge in the prevention of both, onset and progression of
102 CVD as well as of related complications [10].

103 Considering the multiplicity of risk pathophysiology, an integrative approach is needed to
104 identify novel and to validate established CV and metabolic biomarkers for their scientific and
105 clinical utility. Practical biomarkers are required to facilitate (1) the understanding of
106 underlying mechanisms of disease development, (2) the detection of potential targets for
107 specific preventive therapies, and finally (3) the precise estimation of individual risk. For this
108 purpose, there is an unmet need for a cohort studies recruiting individuals at risk for CVD
109 and/or MD well before clinical manifestation of the diseases.

110 In the BioPersMed cohort (Biomarkers of Personalized Medicine), we enrolled community
111 dwelling and asymptomatic individuals from the regional communities who were at risk for
112 CVD or MD in order to evaluate the predictive value of various traditional and novel
113 biomarkers and to observe disease courses starting in the preclinical, asymptomatic phase.
114 The latter shed light on different pathways of CVD and MD development by use of cutting
115 edge laboratory measurements, advanced imaging techniques, comprehensive genetic
116 investigations, and state of the art functional tests. The BioPersMed cohort is located at the
117 Medical University of Graz (Austria) in a dedicated clinical outpatient research centre and
118 biobank (www.biobank.medunigraz.at). The aim of the study is to evaluate large-scale

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3 119 screening tools for the improvement of (1) cardiovascular and metabolic risk stratification, (2)
4 120 early diagnosis of CVD and/or MD (3), individual prediction of clinical outcomes, (4) and long-
5 121 term monitoring of risk and/or early CV and/or metabolic changes in an apparently healthy
6 122 but representative at-risk population at high CV and/or metabolic risk in a prospective
7 123 manner. Ultimately, the data obtained from this cohort aims to facilitate the implementation
8 124 of risk-adapted personalized interventions in both primary and secondary prevention of CVD
9 125 and MD.
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Cohort description

18 127
19 128 The BioPersMed project is designed as a single-centre, prospective, observational study. Only
20 129 asymptomatic subjects without diagnosed CVD but with at least one traditional CV risk factor
21 130 were eligible to participate. According to the published European Guidelines on cardiovascular
22 131 disease prevention in clinical practice, traditional CV risk factors besides of age and gender
23 132 comprise (1) smoking, (2) elevated total cholesterol levels, and (3) arterial hypertension [11].
24 133 Moreover, sedentary lifestyle, obesity, social environment, type 1 diabetes mellitus (T1DM)
25 134 or T2DM, low HDL cholesterol, increased triglyceride levels, elevated fibrinogen,
26 135 apolipoprotein B (apoB), lipoprotein(a), familial hypercholesterinaemia, increased high
27 136 sensitivity (hs)CRP, preclinical evidence of atherosclerosis and chronic kidney disease
28 137 [glomerular filtration rate (eGFR) ≤ 60 mL/min/1.73 m²] were regarded as additional potential
29 138 CV and MD risk factors. From October 2010 (first patient in) to February 2016 (last patient in),
30 139 we enrolled a total of 1022 community dwelling adult men and women who live in the greater
31 140 Graz area via an established recruitment network, consisting of general practitioners,
32 141 peripheral hospitals, and in most cases through the outpatient clinics of the Departments of
33 142 Cardiology as well as Endocrinology and Diabetology, respectively.
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47 143 Subjects presenting with significant non-CVD or who were expected not to be able to
48 144 complete study specific examinations, were excluded from participation. Ethical approval for
49 145 the BioPersMed cohort study has been granted by the Ethics Committee of the Medical
50 146 University of Graz, Austria and is renewed every year (EC Nr. 24-224 ex 11/12). The
51 147 BioPersMed study was conducted in compliance with the laws and guidelines of the Medical
52 148 University of Graz and complies with the Declaration of Helsinki and the Austrian laws [12,13].
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58 149 All participants in the BioPersMed cohort were thoroughly checked for in- and exclusion
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150 criteria before the first phenotyping at baseline examination in order to avoid screening
151 failures.

152 The baseline examinations have been repeated every two years in addition to interim
153 telephone visits, which take place between the on-site visits. A summary of all examinations
154 is shown in figure 1, and a more detailed description can be found in the supplement
155 (Supplemental Tables S1 to S10).

156 According to the presented scheme, participants will be followed for the next decades and
157 clinical outcomes are adjudicated prospectively. A total of 169 (17 %) participants dropped-
158 out for various reasons. Causes for premature unintended termination of the study ranged
159 from a change in the place of residence (n = 5) to limited personal time or lacking will for
160 continuous study visits (n= 136) or new onset of non-CV related diseases (cancer: 4, accident:
161 3, other: 9). Twelve people have died so far (cancer: 7, sepsis: 3, CVD: 2). In summary, 1022
162 persons were included in the baseline examination and 799 persons attended the first follow-
163 up two years after baseline examination. With September 1st, 2021, 628 persons have
164 completed the second follow-up at four years after baseline visit, 531 persons have completed
165 the third follow-up at six years after the baseline visit, and 225 persons have completed the
166 follow-up of eight years after the baseline visit. A small number of participants skipped one
167 follow-up but decided to continue to participate in the study. This issue explains the
168 discrepancy between the number of drop-outs and the number of missing follow-up visits. A
169 detailed overview of the recruitment process and study protocol is presented in figure 2.

170 At baseline and at regular two-years-follow-ups, an in-depth diagnostic CV and MD work-up
171 was carried out, laying emphasis on standardization and reproducibility of history taking,
172 questionnaires for health, psychological and sleep issues, physical examination, ECG,
173 laboratory/blood sampling with biobanking, exercise testing (6min walking test, grip strength,
174 spiroergometry), echocardiographic analysis of cardiac structure and function, pulmonary
175 function testing, carotid intima/media-thickness measurement, pulse-wave analysis, and
176 ophthalmologic examinations as well as body composition, bone density including bone and
177 hormonal biomarkers, and oral glucose tolerance testing. The number of examinations
178 performed at each visit increased over time, due to additional new diagnostic tools (e.g., non-
179 mydriatic funduscopy). A detailed description of all methods used, as well as a concise
180 overview of the assessed data can be found in the supplementary file. Statistics have been
181 calculated using RStudio Version 1.2.5033 (RStudio Inc., United States of America) [14].

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3 182 Besides the already included packages others, namely “hmisc”(Version : 4.6), “ggplot2”(3.3),
4
5 183 “tidyr”(1.12), “readxl”(1.3), “MatchIt”(4.1), “data.table”(1.13), “dplyr”(1.0) and
6
7 184 “lubridate”(1.7), were needed for structuring of data, analysis and visual representation.
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9 185 Normal distribution of data was tested and, if positive, Pearson-correlation calculated. In case
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11 186 of a violation of normal distribution, a non-parametric equivalent was used. A description of
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13 187 the data is given in the corresponding tables. GCP trained and authorized independent
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15 188 persons from the Clinical Trials Unit (CTU) at the Medical University Graz have carefully
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17 189 evaluated and monitored the data and relevant procedures around the cohort study. Based
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19 190 on the steering committees agreement, risk based monitoring of 100 % of the baseline dataset
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21 191 and 80 % of the follow-up dataset was performed. This was done to generate an adequate
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23 192 security of the structure and verification of the assessed data.

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25 194 Biological and technical outlier were manually identified and corrected. This study conforms
26
27 195 to the STROBE-ME recommendations for reporting cohort studies [15].
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30 197 31 32 198 **Health status**

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34 199 At each visit, a careful assessment of the participants’ health status was performed. Data
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36 200 collection documents anthropometric, biochemical, metabolic, hormonal, dietary, physical
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38 201 activity, socioeconomic, medical, and other variables.
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41 203 **Questionnaires**

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43 204 Questionnaires at all clinical visits include the Short-Form 36 Health Survey (SF-36) and the
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45 205 Hospital Anxiety and Depression Scale (HADS), as well as other questionnaires to assess
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47 206 depressive or disease-related symptoms and sleep qualities[16–21]. Raw data of these
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49 207 questionnaires have been collected and are available for further analysis.
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52 209 **Echocardiography**

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54 210 Cardiac chamber geometry and function were assessed via state-of-the-art transthoracic
55
56 211 echocardiography. 2D-, Doppler- and M-mode echocardiography were performed following
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58 212 standardized protocols according to current guidelines and are digitally archived.
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214 **Physical functioning and exercise capacity spiroergometry**

215 For the assessment and interaction of vascular disease, cardiovascular risk and biomarkers, an
216 assessment of physical functioning/exercise capacity was essential. Complementary to
217 questionnaires on physical activity, physical fitness and symptom-limited cardiopulmonary
218 exercise capacity was assessed by spiroergometry, six-minute walk test [22], and handgrip
219 measurements [23,24].

221 **Pulse wave analysis and ECG**

222 Pulse wave analysis (PWA) and pulse wave velocity (PWV) measurements were performed
223 according to the expert consensus published by Laurent et al [25]. Additionally, a 12-lead ECG
224 was performed in every patient.

226 **Ultrasound of the abdomen**

227 Routine abdominal ultrasound examinations with special attention to liver and spleen
228 characteristics (dimensions, texture, abnormalities) were carried out at each visit.

230 **Carotid ultrasound examinations**

231 Carotid ultrasound examinations were performed in supine position on the right and left side.
232 The intima/media thickness of the left and right common carotid artery is measured [26].

234 **Bone density and body composition measurements**

235 Regular DXA (dual energy X-ray absorptiometry) measurements of bone density at the spine,
236 hip and whole-body density include body composition and trabecular bone score (TBS) [27].

238 **Laboratory and functional metabolic measurements**

239 Sample acquisition includes serum, plasma, saliva and urinary as well as stool samples. Routine
240 tests include liver and kidney function and electrolytes, blood counts, hormonal and metabolic
241 data including lipid profiles and urinary analyses. A standardized oral glucose tolerance tests
242 (oGTT, including insulin and c-peptide) was performed. These materials were collected at each
243 visit for immediate analysis as well as biobanking (including samples of function tests).

245 **Biobanking**

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246 Serum, EDTA and citrate plasma, whole blood and cell pellets, spot urine, and saliva were
247 collected at each study visit. They were immediately aliquoted and stored at the biobank of
248 the Medical University of Graz (Austria) at – 80 °C [28]. Biobanking guarantees an accurate
249 description of the sample collection and sample handling according to the STROBE-ME
250 recommendations [15]. Biospecimen-derived measurements adhere to the European
251 guidelines [29].

252

253

254 **Autoantibody phenotyping**

255 Routine measurement of thyroid autoantibodies and autoimmune parameters for celiac
256 disease were performed in all participants. Further exploratory autoantibody detection
257 (endocrine receptors, various epitopes) was done using newly developed luciferase-based
258 fusion protein assays together with the Charité – Universitätsmedizin Berlin, Germany [30].

259

260 **Metabolic phenotyping by Nuclear Magnetic Resonance (NMR)**

261 (Un)targeted metabolomic analysis of 1012 baseline serum and urine samples was performed
262 by NMR spectroscopy and state-of-the art chemoinformatics [31–33]. Metabolite and
263 lipoprotein concentrations were determined using Topspin and the Pre-Clinical Screening and
264 In Vitro Diagnostics Research (IVDr) package of Bruker [34–36].

265

266 **Genome-wide characterisation of cohort**

267 A whole genome global screening array (GSA, Illumina bead chip (Infinium Global Screening
268 Array-24 V2; Illumina Inc, USA)) with nearly 700k specific single nucleotide polymorphisms
269 (SNPs, Illumina, in cooperation with the Human Genotyping Facility at the Erasmus University
270 Rotterdam, NL) is available from all cohort participants.

271

272 **Optometric phenotyping**

273 In addition, a large part of the cohort has been assessed by non-mydratic retinal photography
274 since 2015. Ophthalmologic examinations comprised undiluted fundus photography and
275 optical coherence tomography including the assessment of retinal vessel diameters.

276

277 **Data monitoring and quality assurance**

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278 All incoming data are checked by the study staff for completeness and plausibility and are
279 entered into an electronic data capture system (OpenClinica; www.openclinica.com),
280 specifically adapted for this project. Additional validation processes such as cross-validation
281 with i.e. external independent validation in samples were regularly performed. External
282 monitoring by certified clinical monitors has been done in 100 % of the baseline study records.
283 Adequate external monitoring of the follow-up data is regularly ongoing.

285 **Patient and Public Involvement**

286 No patient involved.

287 **Findings to date**

288 A descriptive overview of the cohort including the cardiovascular, endocrine, and metabolic
289 risk profile is given in table 1. More female (55 %) than male (45 %) persons were included in
290 the study. The mean age of the participants is 57 ± 8 years and the mean BMI is 26.5 ± 4.5
291 kg/m^2 . A majority of 59 % of the examined persons has a BMI greater than 25 kg/m^2 . Although
292 asymptomatic, only 26 % of the study population is considered to be at low cardiovascular risk
293 according to the Framingham “Systematic COronary Risk Evaluation” (SCORE) [37], while 38
294 % show intermediate risk, 26 % show high risk, and 1 % show very high risk. Due to some
295 missing biomarkers at the baseline visit, we were not able to calculate the cardiovascular risk
296 in 9 % of the study population using the Framingham Score. The most common risk factor is
297 hyperlipidaemia (42 % of the population), followed by arterial hypertension (33 % of the
298 population), diabetes mellitus type 2 (T2DM, 18 % of the population), and active smoking (15
299 % of the population). A rather high number, nearly a third of the study population, has been
300 identified as pre-diabetics based on oral glucose tolerance test (oGTT) data.

301 The purpose of this manuscript is to describe the study cohort and to give an overview of the
302 baseline characteristics. These are reported in detail including additional information on used
303 materials and methods in the supplementary file (Tables S1 – S10). The central figure 3 shows
304 an R-correlation plot of most measured biomarkers at baseline. In this plot, biomarkers are
305 not grouped (for example in organ systems); instead, clusters of high correlations were formed
306 (blue indicates a positive correlation, red indicates a negative correlation, and white indicates
307 no correlation) [38]. Thereby, associations between different biomarkers of different organ
308 systems can be revealed which may further serve as a basis for a multi-disciplinary in-depth
309 analysis.

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3 310 Such analyses already identified the so far unknown correlation between IGF1 receptor auto-
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5 311 antibodies with body composition and height as presented at the European Congress of
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7 312 Endocrinology in Barcelona, Spain 2018 [39]. Another preliminary finding revealed a
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9 313 correlation between diabetes status and echocardiographic parameters of the diastolic heart
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11 314 function as presented at the Congress of the European Association for the Study of Diabetes
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13 315 in Barcelona, Spain 2019 [40]. These observations were in line with previous findings of
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15 316 another research group [41]. Furthermore, genetic data from the genome-wide association
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17 317 study and optometric phenotyping of the BioPersMed cohort were already used as control
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19 318 data for a large keratoconus genomic study in cooperation with researchers from the United
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21 319 Kingdom and the Netherlands [42], for the analysis of allelic determinants with a reported
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23 320 association to 25(OH)D levels and their influence on vitamin D [43], and to identify novel
24
25 321 biomarkers for non-alcoholic fatty liver disease [44] in cooperation with local researchers.
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27 322 Data on bone morphology from the BioPersMed cohort were used to link Sarcopenia with
28
29 323 increased risk of falls, osteoporosis and mortality[45]. In addition, miRNA profiles were linked
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31 324 to Hashimoto's Disease and Thyroid Antibodies[46].
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Strengths and limitations

36 328 The main strength of the BioPersMed cohort is the joint evaluation of both cardiovascular and
37
38 329 metabolic phenotyping including a broad spectrum of diagnostic, imaging, and functional
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40 330 tools. This assures comprehensive biomedical and scientific dimensions of the project within
41
42 331 high-end diagnostic and analytical parameters and biomarkers. Second, biobanking with a
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44 332 large number of samples including serum/plasma and blood cells, urine, saliva and stool at
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46 333 each visit that have been aliquoted and stored upon high quality certification rules enabling a
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48 334 prospective view on candidate biomarker in the context of a large longitudinal cohort, where
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50 335 specific approaches can be predefined [47,48]. In addition, a specifically adapted large
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52 336 electronic database (OpenClinica; www.openclinica.com) assures the quality of data entry and
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54 337 delivery as well as the validity and reliability of biomarkers analyses in the BioPersMed cohort
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56 338 together with a continuous data monitoring.
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58 339 A potential weakness of this study is the wide time range of a prolonged recruitment due to
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60 340 logistic reasons between 2010 and 2016. This results in a prolonged follow-up period of study
341 participants between 4 and 10 years. Second, after a thorough standardized baseline

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3 342 phenotyping, of the BioPersMed cohort this phenotyping has been expanded by additional
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5 343 diagnostic parameters at either later points in time (e.g. non-mydratic fundoscopy) or in
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7 344 subpopulations. Although some of these biomarkers are not available for the complete
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9 345 duration of the entire cohort, cross-sectional analysis of a considerable number of participants
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11 346 can already be performed with these data sets and will be available for longitudinal
12
13 347 comparison of follow-up visits thereafter.
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349 **Collaboration**

16 349
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18 350 The design of the BioPersMed study, data management, biobanking and data analyses are
19
20 351 compliant to the STROBE, STROBE-ME and STREGA recommendations. Collaboration in data
21
22 352 analysis and publications will be welcome through specific research proposals sent to the
23
24 353 BioPersMed investigators listed as corresponding authors of this manuscript. If desired,
25
26 354 retrospective analyses can be performed because all data are recorded as raw data and more
27
28 355 than 300,000 samples of blood, serum, and urine are stored in the biobank.
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31 **Further Details**

32 **Data sharing statement**

33 358
34
35 359 Data are available upon request. To get access, a proposal must be submitted to the
36
37 360 investigators listed as the corresponding authors of this manuscript
38
39 361 (*barbara.obermayer@medunigraz.at* and *albrecht.schmidt@medunigraz.at*). Additional
40
41 362 information can be obtained via the Research Management of the Medical University of Graz
42
43 363 (*tanja.ball@medunigraz.at*)
44
45 364

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373 Cardiovascular Risk Assessment"; THEME HEALTH.2011.2.4.2-2; Grant agreement no:

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376 Graz; Austrian Infrastructure Program 2016/2017, the Styrian Government (Zukunftsfonds).

377

378 **Contributorship statement**

379 BP, TP, AS, and BOP designed and supervised the study. AS and BOP administrated the study.
380 CH, EK, NV, BOP, and AS wrote the manuscript. EK, CC, IM, MUM, TG, BH, AL, CR, NV, KA, MW,
381 EPK, NT, GS, and AS investigated participants and collected data. CH, EK calculated the
382 statistics performed principal data analysis. AT, NS, NV, TM, AS, AW, AZ, TK, and RS
383 contributed with data analysis and the allocation of resources. All authors reviewed the
384 manuscript

385

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392 been provided by Biobank Graz of the Medical University of Graz, Austria. Cohort 5001_15,
393 BioPersMed.

394

395 **Conflict of interest:**

396 The authors declare no conflict of interest.

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3 398 **Figures**

4
5 399 **Figure 1**

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11 402 **Figure 1**

12 403 Illustration of the comprehensive phenotyping and biosampling of the BioPersMed Cohort.

13 404 Follow-up visits are performed according to a tight pre-planned schedule, including reminder-

14 405 phone calls by study nurses with phenotyping and biosampling every second year and a follow-

15 406 up telephone visit in the years between. At baseline, every participant received a patient's

16 407 diary for documentation of medical events (source data).

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3 409 **Figure 2**

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10 413 **Figure 2**

11 414 The recruitment status over time and the follow-up of the participants' phenotyping in the

12 415 BioPersMed Cohort. A: Absolute number of participants who have completed various on-site

13 416 follow-ups until 1st September 2021. B: Timeline of recruitment and follow-up processes.

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3 418 **Figure 3**

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9 421 **Figure 3**

10 422 R-plot of the analysed data set. Potential biomarkers are not grouped; instead, random
11 423 clusters of correlations are formed. Blue zones indicate positive correlations, red zones
12 424 indicate negative correlations. As an example, zones of strong correlations are zoomed and
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14 425 potential biomarkers are depicted.
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427 **Tables**428 **Table 1**

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	Number (N total =1022)	%
Sex		
Men	455	45
Women	567	55
Smoking		
Active smoker	154	15
Former smoker	326	32
Non-smoker	506	49
Unknown	36	4
Framingham SCORE*		
Low Risk (< 3 %)	263	26
Intermediate Risk (3 – 4 %)	390	38
High Risk (5 – 9 %)	267	26
Very High Risk (≥ 10 %)	9	1
Unknown	93	9
Diabetes status**		
NG	390	38
PreD	315	31
T2DM	181	18
unknown	136	13
Medical history***		
Arterial Hypertension	341	33
Hyperlipidemia	434	42
Stroke / TIA	7	1
Antihypertensive drugs per patient		
No antihypertensive drug	670	65
One antihypertensive drug	154	15
Two antihypertensive drugs	118	12
More than two antihypertensive drugs	80	8
Lipid-lowering drugs per patient		
No lipid-lowering drug	876	85
One lipid-lowering drug	139	14
More than one lipid-lowering drugs	7	1
Antidiabetic drugs per patient		
No antidiabetic drug	844	82
Dietetic treatment	119	12
One antidiabetic drug	54	5

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More than one antidiabetic drugs	5	1
Age		
< 55 years	437	43
55 – 65 years	382	38
66 – 75 years	178	17
> 75 years	25	2
Body mass index		
< 18.5 kg/m ²	6	1
18.5 – 25.0 kg/m ²	408	40
> 25.0 kg/m ²	608	59

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431 Health status - abbreviations

432 NG = normoglycemia; T2DM = type 2 diabetes mellitus; PreD = prediabetes ; TIA = transient ischemic attack ; *
 433 whenever present, a direct categorization due to comorbidities was performed; **Diabetes status based on oGTT
 434 results; *** Hyperlipidemia was assumed based on total cholesterol > 200 mg/dL or the use of a lipid-lowering
 435 drug

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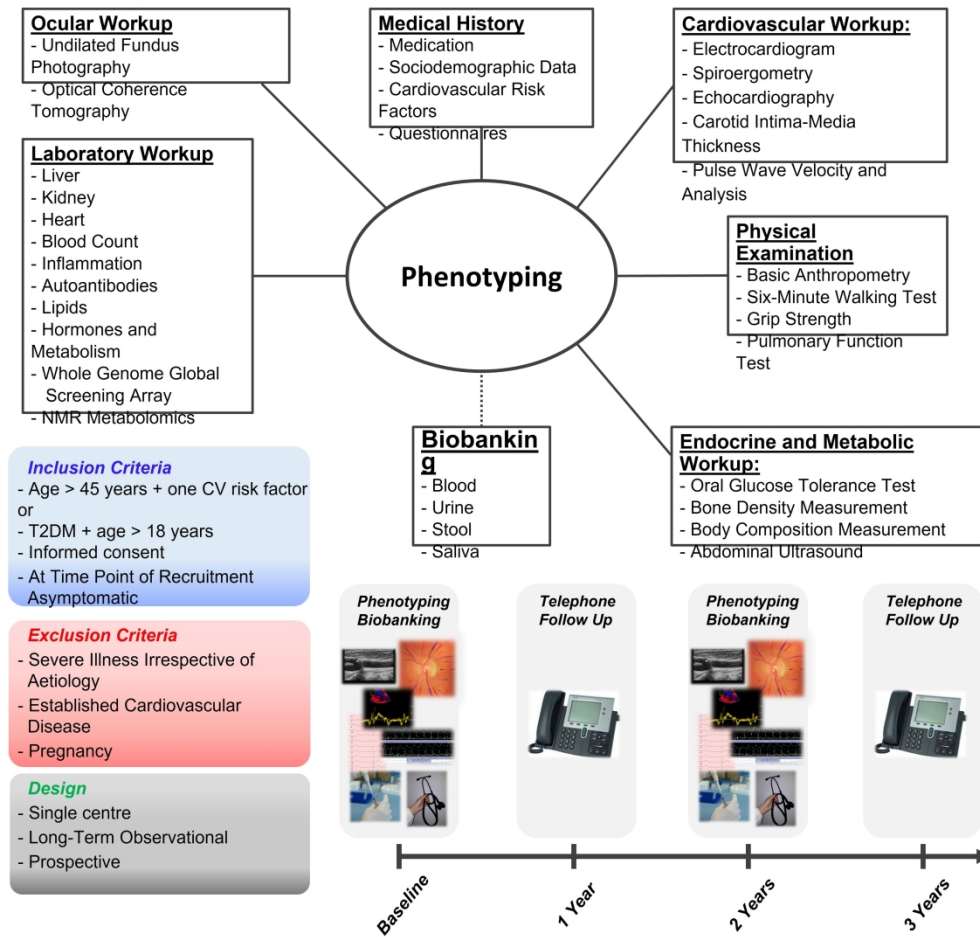
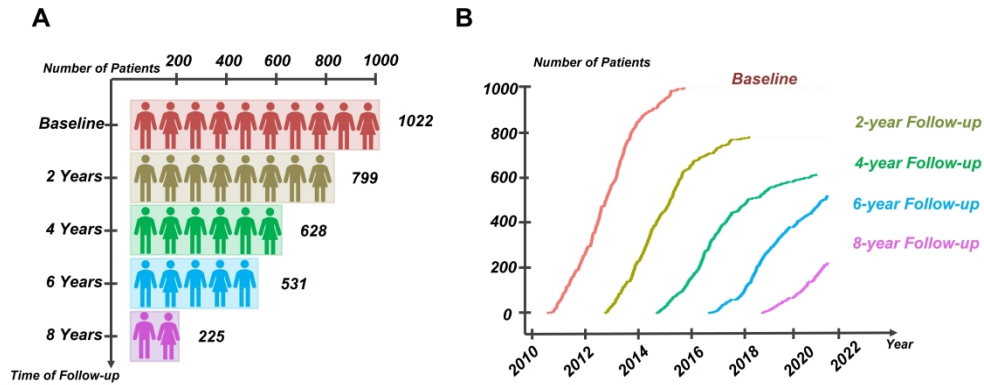


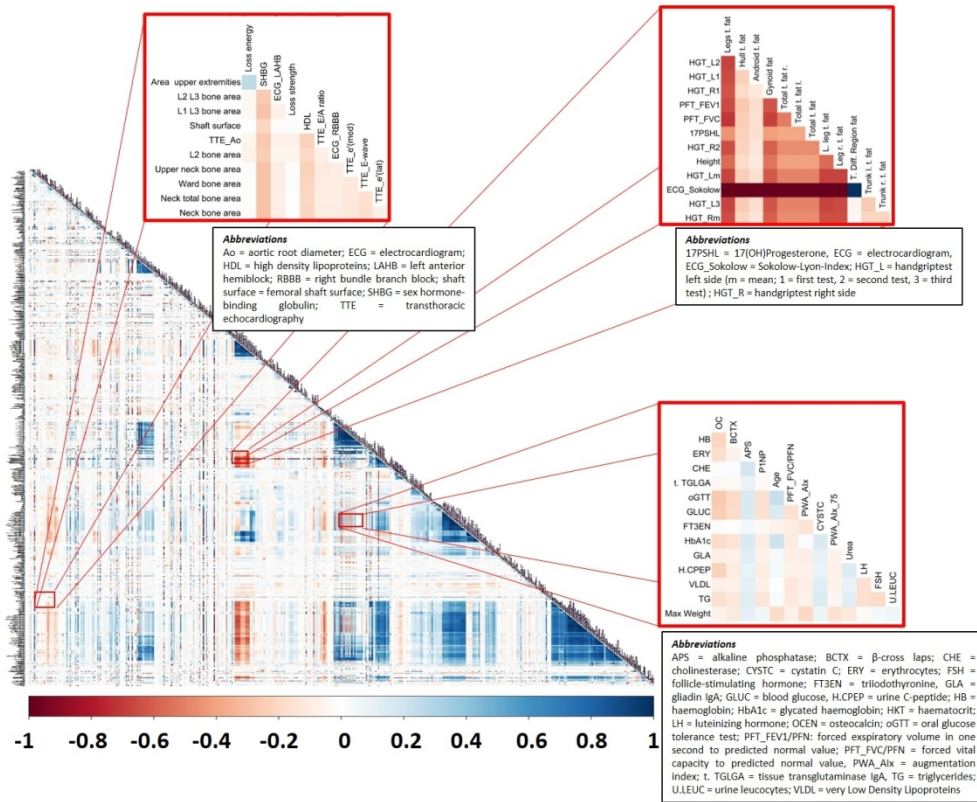
Illustration of the comprehensive phenotyping and biosampling of the BioPersMed Cohort. Follow-up visits are performed according to a tight pre-planned schedule, including reminder-phone calls by study nurses with phenotyping and biosampling every second year and a follow-up telephone visit in the years between. At baseline, every participant received a patient’s diary for documentation of medical events (source data).

320x304mm (300 x 300 DPI)



The recruitment status over time and the follow-up of the participants' phenotyping in the BioPersMed Cohort. A: Absolute number of participants who have completed various on-site follow-ups until 1st September 2021. B: Timeline of recruitment and follow-up processes.

441x192mm (330 x 330 DPI)



R-plot of the analysed data set. Biomarkers are not grouped; instead, random clusters of correlations are formed. Blue zones indicate positive correlations, red zones indicate negative correlations. As an example, zones of strong correlations are zoomed and biomarkers are depicted.

256x211mm (150 x 150 DPI)

BioPersMed Concept Paper

Supplements – Methods and baseline characteristics

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1. Echocardiography

For all cardiac ultrasound examinations, a GE vivid E9 device (GE Healthcare, United Kingdom) with a phased array transducer (GE M5s) was used. During the echocardiographic examination, patients were positioned in the left lateral decubitus position. An ECG tracing was recorded during the whole examination. Loops of 3 cardiac cycles of the parasternal long axis view, the parasternal short axis view (on the level of the aortic valve, the mitral valve, the papillary muscles, and the apex), the apical four chamber view, the apical two chamber view, the apical three chamber view, the apical five chamber view, and the subcostal view were recorded. Additionally, colour Doppler loops of all mentioned echocardiographic angulations were recorded for the semi-quantitative assessment of a regurgitation or stenosis. Doppler measurements of the transmitral flow, the transaortic flow, the tricuspid regurgitation flow, and the pulmonic venous flow were recorded as images. Cardiac chambers were quantified by the measurement with the M-Mode technique or as a measurement within the two-dimensional image. M-mode and Doppler images were recorded at a sweep speed of 100 mm/s. The colour Doppler measurements were recorded with a Nyquist limit of 65 ± 5.5 cm/s. For the calculation of the left ventricular mass, the modified Devereux formula [1] was used, for the calculation of the left atrial volume, the biplane formula was used. The body surface area was calculated from height and weight using the DuBois formula. Table S1 gives an overview of all assessed parameters of the baseline-examination.

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Table S1: Echocardiographic parameters of the baseline cohort (N=1022)			
	Method	Mean \pm SD	Unit
Left Ventricle			
IVS end diastolic - PLAX	M-Mode	10.5 \pm 1.9	mm
PW end diastolic - PLAX	M-Mode	10.6 \pm 2.0	mm
LVD end diastolic - PLAX	M-Mode	47.4 \pm 5.9	mm
LVM	Calculated (modified Devereux formula)	184.5 \pm 61.3	g
LVMi	Calculated (LVM indexed to BSA)	96.8 \pm 26.2	g/m ²
LVD end systolic PLAX	M-Mode	29.8 \pm 4.8	mm
LVD apical major axis end diastolic - A4CH	2D	76.1 \pm 8.1	mm
LVD apical minor axis end diastolic - A4CH	2D	43.6 \pm 5.5	mm
LVV end diastolic - A4CH + A2CH	2D (Simpson's biplane method)	91.4 \pm 26.9	mL
LVV end systolic - A4CH + A2CH	2D (Simpson's biplane method)	34.0 \pm 13.1	mL
EF	Calculated (Simpson's law)	63.8 \pm 5.8	%
MAPSE medial annulus - A4CH	M-Mode	0.92 \pm 0.16	cm
MAPSE lateral annulus - A4 CH	M-Mode	1.54 \pm 0.26	cm
Right Ventricle			
TAPSE - A4CH	M-Mode	2.31 \pm 0.09	cm
Left Atrium			
LA minor axis end systolic - PLAX	PLAX	38.4 \pm 5.7	mm
LA major axis end systolic - A4CH	2D	48.0 \pm 6.0	mm
LAA end systolic - A4CH	2D	17.6 \pm 3.6	cm ²
LAA end systolic - A2CH	2D	17.9 \pm 3.6	cm ²
LAV end systolic - A4CH + A2CH	Calculated (biplane method)	56.1 \pm 16.6	mL
LAVI end systolic - A4CH + A2CH	Calculated (LAV indexed to BSA)	29.8 \pm 7.9	mL/m ²
Doppler Measurements			
AV-Vmax	CWD	1.2 \pm 0.3	m/s
Peak E-wave velocity	PWD of transmitral flow	70.0 \pm 16.1	cm/s
Peak A-wave velocity	PWD of transmitral flow	67.2 \pm 18.1	cm/s
Mitral valve deceleration time	PWD of transmitral flow	221 \pm 56	ms
E/A ratio	PWD of transmitral flow	1.10 \pm 0.38	
Duration A-wave	PWD of transmitral flow	132 \pm 37	ms
Duration IVRT	PWD of transmitral flow	115 \pm 20	ms
Duration aortic flow	PWD of transmitral flow	276 \pm 37	ms
Duration E-wave to A-wave	PWD of transmitral flow	447 \pm 48	ms
Peak E-wave velocity (<i>valsalva</i>)	PWD of transmitral flow under valsalva	62.2 \pm 16.0	m/s
Peak A-wave velocity (<i>valsalva</i>)	PWD of transmitral flow under valsalva	63.3 \pm 18.6	m/s
Mitral valve deceleration time (<i>valsalva</i>)	PWD of transmitral flow under valsalva	241 \pm 67	ms
e' velocity (medial annulus)	TDI-PWD	7.6 \pm 2.2	cm/s
e' velocity (lateral annulus)	TDI-PWD	10.4 \pm 3.0	cm/s
E/e' mean	Calculated	8.2 \pm 2.4	
Tricuspid regurgitation systolic jet velocity	CWD	2.25 \pm 0.27	m/s
Pulmonary vein systolic velocity	PWD of pulmonary venous flow	56.7 \pm 12.5	cm/s
Pulmonary vein diastolic velocity	PWD of pulmonary venous flow	42.6 \pm 10.1	cm/s
Pulmonary vein S/D ratio	PWD of pulmonary venous flow	1.37 \pm 0.38	
Flow propagation time	CD M-Mode of transmitral flow	68.6 \pm 31.0	cm/s
Valves			
Semiquantitative evaluation of regurgitation	CD		

Echocardiographic angulations and Doppler measurements - abbreviations:

PLAX = parasternal long axis view; A4CH = apical four chamber view; A2CH = apical two chamber view; CWD = continuous wave Doppler; PWD = pulsed wave Doppler; TDI = tissue Doppler imaging; CD = colour Doppler

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Cardiac structures - abbreviations

IVS = interventricular septum; PW = posterior wall; LVD = left ventricular diameter; LVM = left ventricular mass; LVMI = left ventricular mass index; LVV = left ventricular volume; EF = ejection fraction; MAPSE = mitral annulus plane systolic excursion; TAPSE= tricuspid annulus plane systolic excursion; LA = left atrium; LAA = left atrial area; LAV = left atrial volume; LAVI = left atrial volume index; IVRT = isovolumetric relaxation time

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2. Physical functioning and exercise capacity spiroergometry

The whole study cohort underwent several examinations assessing the cardiac, pulmonary, and exercise capacity. Firstly, a blood pressure measurement with the Boso-medicus uno (Bosch + Sohn GmbH, Germany) was performed in the seating position after 10 minutes of rest. During the diagnostic workup, a spirometry was performed ahead of ergometry. Symptom limited cardiopulmonary exercise testing on a bicycle ergometer starts at a workload of 20W, followed by a stepwise 20W increment every 2 min. Criteria for discontinuation of the exercise test are defined as recommended by the European Society of Cardiology [2]. A standard 12-lead ECG continuously monitors heart rate, ST-segment changes, and arrhythmias. Blood pressure is recorded at rest and then every 2 min. Ventilatory exchange, oxygen uptake (VO_2), and other cardiopulmonary variables are acquired by averaging breath-by-breath measurements over 10 s intervals. Peak heart rate and workload is recorded immediately upon the end of exercise. Peak VO_2 is defined as the maximum value of the last three 10 s averages during exercise and anaerobic threshold is detected using the V-Slope method [3], based on standard operating procedures for spiroergometry. Finally, participants reported the degree of exhaustion expressed as the level of shortness of breath according to a Borg category-ratio scale (0 – 10 points)[4].

After a period of at least 15 minutes rest, a six-minute walk test was performed. Initially, resting blood pressure was assessed; then patients were instructed to walk for six minutes as far as possible without running or jogging. The Borg category-ratio scale (0 – 10 points) was assessed after the six-minute walking test too.

Finally, a handgrip test using a Jamar® Hydraulic Hand Dynamometer (Performance Health UK, United Kingdom) was performed to measure grip strength of both hands. The participant was tested in the seated position with elbow flexed at 90 degrees without touching the trunk.

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The handle position was adjusted to fit the participant's hand. The participant was asked to squeeze the dynamometer at maximal effort for three trials, with a 30-second break between each trial. The average of three trials was calculated for data analysis. Table S2 gives an overview on the single assessed physical functioning and exercise capacity related biomarkers.

Table S2: Baseline parameters of spirometry, spiroergometry, 6-minute walking test and hand-grip strength test			
	Male	Female	Unit
Blood pressure measurement (N=1022)			
SBP – seating position	141 ± 19	133 ± 20	mmHg
DBP – seating position	88 ± 11	85 ± 11	mmHg
Spirometry (N=1022)			
FVC	4.1 ± 0.9	3.0 ± 0.6	L
PNV (FVC)	4.6 ± 0.5	3.1 ± 0.4	L
FVC / PNV	90.3 ± 16.1	97.1 ± 17.9	%
FEV1	3.3 ± 0.8	2.4 ± 0.5	L/s
PNV (FEV1)	3.5 ± 0.4	2.5 ± 0.4	L/s
FEV1 / PNV	93.2 ± 20.0	94.3 ± 19.7	%
FEV1 / FVC	102.3 ± 17.6	102.3 ± 19.8	%
Spiroergometry (N=1022)			
Wmax	158 ± 36	111 ± 23	W
SBP – resting	125 ± 19	118 ± 19	mmHg
DBP – resting	83 ± 12	78 ± 11	mmHg
SBP - maximal	206 ± 23	189 ± 25	mmHg
DBP - maximal	90 ± 16	89 ± 18	mmHg
HR – resting	77 ± 13	79 ± 13	/min
HR - maximal	143 ± 20	147 ± 19	/min
Peak VO ₂	23.9 ± 7.0	20.5 ± 6.4	mL/kg/min
Anaerobic threshold	107 ± 41	79 ± 25	W
Borg-CR10	3.9 ± 2.0	3.9 ± 2.0	
Six-minute walk test (N=980)			
SBP – resting	139 ± 18	133 ± 20	mmHg
DBP – resting	88 ± 11	85 ± 11	mmHg
SBP - maximal	147 ± 22	141 ± 22	mmHg
DBP - maximal	89 ± 11	88 ± 12	mmHg
HR – resting	73 ± 14	73 ± 11	/min
HR - maximal	80 ± 16	82 ± 14	/min
Distance walked	521 ± 90	501 ± 91	m
Prematurely terminated	3 / 446	4 / 559	N / total
Borg-CR10	0.4 ± 0.8	0.6 ± 1.2	
Hand-grip test (N=680)			
Right arm	96 ± 20	59 ± 13	lbs
Left arm	90 ± 20	54 ± 13	lbs

spiroergometry, six-minute walk test & hand-grip test - abbreviations

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FVC = forced vital capacity; PNV = predicted normal value; FEV1 = forced expiratory volume (1 second); Wmax = maximal workload; SBP = systolic blood pressure; DBP = diastolic blood pressure; HR = heart rate; VO₂ = oxygen uptake; Borg-CR10 = Borg category-ratio scale

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3. Pulse wave analysis and ECG

Measurements of pulse wave analysis (PWA) and pulse wave velocities (PWV) were performed with the SphygmoCor® device (Atcor Medical, Australia). The measurement takes place in supine position after a minimum of 10 to 15 min rest in a quiet, temperature-controlled room after measuring the blood pressure on the right and left side [5]. Central BP measurements were recorded noninvasively by applanation tonometry.

PWV (carotid–femoral), which reflects arterial stiffness, were measured by using the foot-to-foot velocity method. The waveforms were recorded transcutaneously at the right common carotid artery and the right femoral artery. In addition, the augmentation index was determined by applanation tonometry on the central pressure waves measured in the right radial artery according to previous recommendations [6]. The SphygmoCor device quality index (QI), which represents waveform reproducibility, was checked after each pulse wave analysis (PWA) and the measurements are repeated in case of an $QI < 80$. As an instant quality check for the measurement of the PWV, the time difference between the ECG-signal and the signal from the recording sites was suggested to have a $SD \leq 10\%$ of the mean value. All measurements of PWA and PWV were performed in duplicate.

Briefly after the pulse wave analysis, a 12-lead resting ECG was performed. An overview of the ECG and PWA related biomarkers is shown in table S3.

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Table S3: Baseline parameters of 12-lead ECG and pulse wave velocity/analysis		
	Mean \pm SD	Unit
12-lead ECG (N=1019)		
HR	64.4 \pm 11.1	/min
PR-Interval	159.3 \pm 24.9	ms
QRS-duration	95.8 \pm 15.9	ms
QTc-time	418.6 \pm 27.6	ms
PWA (N=993)		
SBP	126 \pm 18	mmHg
DBP	86 \pm 11	mmHg
AP	12.2 \pm 7.6	mmHg
Alx	28.2 \pm 11.4	%
Alx(HR75)	23.9 \pm 11.3	%
PP	39.9 \pm 13.1	mmHg
PWV (N=963)		
Carotid-femoral PWV	7.4 \pm 2.1	m/s

12-lead-ECG and pulse wave analysis – abbreviations

HR = heart rate; SBP = systolic blood pressure; DBP = diastolic blood pressure; AP = augmentation pressure; Alx = augmentation index; Alx(HR75) = augmentation index normalized to the heart rate of 75 beats per minute; PP = pulse pressure

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4. Ultrasound of abdomen

An ultrasound of the abdomen was performed using the GE vivid E9 ultrasound device with a curved array transducer (GE CI5). The dimensions of the spleen were assessed, and the echogenicity of the liver was evaluated in order to identify signs of liver-steatosis. The findings of the abdominal ultrasound are summarized in table S4.

	Number (N total = 982)	%
Ultrasound of abdomen		
Normal	626	64
Blunted liver edge	32	3
Irregular liver surface	2	< 1
Ascites	1	< 1
Signs of steatosis	321	33
Dimension of spleen	Mean \pm SD	Unit
Longitudinal	101.8 \pm 14.4	mm
Transversal	36.6 \pm 7.7	mm

5. Carotid ultrasound examinations

Carotid ultrasound examination was performed with a GE vivid E9 ultrasound device with a linear array transducer (GE 9L). Each measurement includes both the near-wall and far-wall data respectively for the right and left carotid arteries [7]. The common carotid intima-media thickness (CIMT) was calculated by an automated measurement program by GE at more than 200 points distal from the bulbous. Sonographers are taking images showing the maximum thickness of a particular site. Detected plaques at specific sites are included in the maximum CIMT measurement. Table S5 gives an overview on the biomarkers of the carotid ultrasound examination.

Table S5: Baseline parameters of carotid ultrasound		
	Mean \pm SD	Unit
CIMT		
Right side	0.73 \pm 0.16	mm
Left side	0.73 \pm 0.17	mm
Presence of calcified plaques or stenosis	Number (N=1021)	%
Right side	117	11
Left side	106	10

Carotid ultrasound – abbreviation:

CIMT = carotid intima-media thickness

6. Bone density and body composition

Bone imaging via 2D bone density measurements and body composition measurements were performed using Lunar iDXA (General Electrics, USA) at each visit including measurements at the lumbar spine, the hip and whole-body sites. Thorough body composition, automated comparison to prior measurements and an additional trabecular bone score (TBS, as a measure of bone texture correlated with bone microarchitecture, software by iDEXA solutions, Switzerland [8]) has been documented. Table S6 shows an overview on the parameters of the bone density measurements.

	Male (N=341)	Female (N=448)	%
Bone mineral density			
normal BMD	325	412	93
Osteopenia or osteoporosis	16	36	7
Body Composition	Mean ± SD	Mean ± SD	Unit
Total BMD	1.28 ± 0.11	1.13 ± 0.12	g/cm ²
Total BMC	3132 ± 392	2293 ± 336	g
Total T-Score	0.78 ± 1.09	0.48 ± 1.14	STD
Total Z-Score	0.63 ± 1.04	1.01 ± 0.94	STD
Total Tissue mass	84397 ± 13301	68502 ± 13681	g
Total Bone mass	3194 ± 243	2180 ± 181	g
L1-L4 BMD	1.25 ± 0.19	1.14 ± 0.17	g/cm ²
L1-L4 BMC	82.2 ± 16.4	62.4 ± 12.5	g
L1-L4 T-Score	0.09 ± 1.52	-0.43 ± 1.36	STD
L1-L4 Z-Score	0.14 ± 1.51	0.29 ± 1.19	STD

Bone density measurements – abbreviations:

BMD = Bone mineral density in [g/cm²], BMC = Bone mineral content [g], L1-L4 = region vertebrae L1-L4, T-Score= standard deviations from mean BMD level of young adults, Z- or T-Score STD = standard deviations from age adjusted mean BMD level

7. Laboratory assessment and measurements

Routine clinical laboratory parameters such as blood count and coagulation, serum electrolytes, kidney and liver function, HbA1c, fasting c-peptide, insulin as well as fasting blood glucose followed by a standardized oral glucose tolerance test (oGTT, at 30, 60 and 120 minutes after 75mg glucose load) with consecutive insulin and c-peptide measurements, high-sensitive c-reactive protein (hsCRP), lipids (total cholesterol, high-density and low-density lipoproteins (HDL and LDL), lipoprotein a (LP(a), triglycerides (TG), high-sensitive troponin T (hsTropT) and cardiac parameters such as creatine kinase (CK, CK-MB), as well as urinary proteins, electrolytes, cell count and c-peptide are measured in all participants at each study visit. The baseline measurements are presented in table S7.

In addition to the oGTT and glucose metabolism monitoring, a large panel of hormonal and metabolic analyses are performed in women as well as in men. These biomarkers include estrogen, testosterone and free testosterone, androstenedione, dehydroepiandrosterone (DHEAS), 17(OH)progesterone, and anti-Müllerian hormone (AMH), saliva hormones (e.g. cortisol), bone metabolism including formation and resorption parameters (osteocalcin, N-terminal propeptide of type 1 procollagen (P1NP), serum-crosslaps (CTX)), calciotropic hormones such as 25(OH)vitamin D and parathyroid hormone (PTH), pituitary hormones (LH, FSH, ACTH) and others.

Routine biochemical parameters (liver, kidney, lipid, electrolyte and other profiles) were determined using a Cobas® Analyzer (Roche Diagnostics, Germany), complete blood counts by Beckmann-Coulter, Germany. Hormonal measurements such as insulin and c-peptide were measured by chemiluminescence on an Advia Centaur system (Siemens Healthcare Diagnostics, Germany). Automated analysers were used to measure hormones: anti-Müllerian hormone (AMH) (Beckmann-Coulter, Germany); testosterone, cortisol, thyrotropin,

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triiodothyronine and thyroxine (Siemens Advia Centaur, Germany); sexual hormone-binding globulin (SHBG) (Roche Diagnostics, Germany); estrogen, luteotropic hormone (LH) and follicle-stimulating hormone (FSH) (Triturus, Biomedical Diagnostics, Antwerp, Belgium); 25(OH)vitamin D and bone metabolism parameters such as bone-specific alkaline phosphatase (bALP), osteocalcin (OC), C-terminal telopeptide (CTX) and procollagen type 1 N-terminal propeptide (P1NP) by iSYS, (IDS, UK, or Roche Diagnostics, Germany, respectively). Endocrine and bone biomarkers are shown in table S8, all of them collected in the morning after an overnight fast.

	Male (N=444)	Female (N=556)	Unit
Liver	Mean ± SD	Mean ± SD	
ALP	93 ± 32	109 ± 11	U/L
ALT	24.7 ± 31.3	16 ± 7.3	U/L
AST	20.9 ± 17.1	18.1 ± 7.4	U/L
CHE	11059 ± 928	11161 ± 1153	U/L
GGT	44.8 ± 54.2	18 ± 20.1	U/L
LDH	161 ± 19	166 ± 19	U/L
Kidney			
Creatinine	0.98 ± 0.14	0.8 ± 0.13	mg/dL
Creatinine/urine	142 ± 35	130 ± 45	mmol/L
Urea	18.8 ± 1.5	18.1 ± 2.2	mg/dL
Uric Acid	6.3 ± 1.2	4.8 ± 1.2	mg/dl
Proteins			
Albumin	4.6 ± 0.3	4.6 ± 0.3	g/dL
CRP	2.3 ± 4.1	2.5 ± 3.7	pg/mL
Ferritin	158 ± 173	123 ± 47	ng/mL
Total Protein	7.4 ± 0.4	7.4 ± 0.4	g/dL
Transferrin	2.5 ± 0.3	2.5 ± 0.3	g/L
Electrolytes & Iron			
Calcium/urine	2.41 ± 1.88	2.21 ± 1.69	mmol/L
Chloride	103 ± 2	104 ± 2	mmol/L
Chloride/urine	149 ± 33	135 ± 41	mmol/L
Iron	132 ± 24	126 ± 22	µg/dL
Magnesium	0.85 ± 0.06	0.85 ± 0.06	mmol/L
Phosphate	2.61 ± 0.75	2.96 ± 0.49	mg/dL
Phosphate/urine	18.6 ± 14.5	15.1 ± 10.8	mg/dl
Potassium	4.1 ± 0.5	4.0 ± 0.3	mmol/L
Total Calcium	2.39 ± 0.09	2.39 ± 0.1	mmol/L
Haematopoiesis			
Haemoglobin	15.1 ± 1.0	13.6 ± 0.9	g/dL
Platelets	174 ± 20	180 ± 18	10 ⁹ /L
TLC	5.77 ± 1.44	5.76 ± 1.76	10 ⁹ /L
Glucose metabolism			
Glucose fasting	99 ± 20	91 ± 17	mg/dL

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Glucose 30 min	158 ± 38	142 ± 36	mg/dL
Glucose 60 min	158 ± 62	128 ± 52	mg/dL
Glucose 120 min	115 ± 57	100 ± 46	mg/dL
Insulin fasting	13 ± 17	10 ± 8	μU/mL
Insulin 30 min	64 ± 54	56 ± 48	μU/mL
Insulin 60 min	94 ± 87	73 ± 68	μU/mL
Insulin 120 min	62 ± 72	51 ± 62	μU/mL
HbA1c	38.6 ± 7.1	37.7 ± 5.8	mmol/mol
Heart			
CK	151 ± 132	130 ± 25	IU
NT-Pro-BNP	79 ± 18	122.66 ± 105.7	pg/mL
PT	105.7 ± 5.0	106.8 ± 5.2	%
Lipid parameters			
Cholesterol	174.8 ± 18.9	181.6 ± 16.6	mg/dL
LDL	130.9 ± 36.5	132.1 ± 33.6	mg/dL
VLDL	19.6 ± 21.9	15.7 ± 2.8	mg/dL
HDL	48.5 ± 16.3	65.1 ± 20.2	mg/dL
LPA	1.37 ± 3.47	1.81 ± 4.16	mg/dL
Triglycerides	144 ± 80	133 ± 26	mg/dL

Laboratory measurements – abbreviations:

ALP = Alkaline Phosphatase; ALT = Alanine Aminotransferase; AST = Aspartate Aminotransferase; CHE = Cholinesterase; CRP = C-Reactive Protein; GGT = Gamma Glutamyl Transferase; LDH = Lactate Dehydrogenase; HB = Haemoglobin; PLT = Platelets; TLC = Total leucocyte count; HbA1c = glycated haemoglobin; CK = Creatine Kinase; NT-Pro-BNP = N-terminal pro brain natriuretic peptide; PT = Prothrombin Time; LDL = Low Density Lipoprotein; VLDL = Very low Density Lipoprotein; HDL = High Density Lipoprotein; LPA = Lipoprotein A

	Male (N=446)	Female (N=557)	Unit
Endocrine parameters			
	Mean ± SD	Mean ± SD	
17(OH)Progesterone	0.9 ± 0.4	0.4 ± 0.3	ng/ml
ACTH	21.3 ± 11.4	14.7 ± 8.1	pg/ml
Aldosterone	8.7 ± 7.6	9.0 ± 8.1	ng/dl
Androstenedione	2.6 ± 1	1.9 ± 0.9	ng/ml
Cortisol	123.2 ± 36.5	111.9 ± 38.13	ng/ml
Cortisol/saliva	7 ± 4.4	6.4 ± 4.2	ng/ml
DHEAS	1.1 ± 0.7	0.7 ± 0.5	μg/ml
Estradiol	58.4 ± 280.8	49.4 ± 63.8	pg/ml
Free Thyroxine	15.1 ± 2.2	15.1 ± 2.7	pmol/l
Free Triiodothyronine	5 ± 0.5	4.6 ± 0.6	pmol/l
FSH	9.34 ± 8.7	67.2 ± 30.9	mU/ml
Insulin	13.4 ± 16.9	9.9 ± 7.7	mU/l
LH	6.0 ± 5.7	14.2 ± 6.4	mU/ml
Progesterone	0.51 ± 1.18	0.9 ± 4.06	ng/ml
Renin	58.3 ± 106.5	30.9 ± 74.4	μU/ml
SHBG	41.1 ± 42.1	119.5 ± 51.8	nmol / L
Total Testosterone	11.0 ± 3.9	1.4 ± 0.7	pg/ml
TSH	2.0 ± 1.1	2.2 ± 6.3	μU/ml
Bone parameters			
25(OH)Vitamin D	32.4 ± 11.3	33.5 ± 11.9	ng/ml
Crosslaps/serum	0.3 ± 0.12	0.37 ± 0.17	ng/ml

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Osteocalcin	18.8 ± 5.6	23.2 ± 8.7	ng/ml
P1NP	40.4 ± 13.6	49.8 ± 20.4	ng/ml

Laboratory measurement: endocrinology & bone metabolism - abbreviations:

ACTH = Adrenocorticotrophic hormone DHEAS = dehydroepiandrosterone sulfate, FSH = Follicle-stimulating hormone, LH = Luteinizing hormone, SHBG = Sex hormone-binding globulin, TSH = Thyroid-stimulating hormone, thyrotropin, P1NP = procollagen type 1 N-terminal propeptide

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8. Autoantibody phenotyping

Data are available for routine thyroid and gluten autoantibodies, further autoimmune parameters for glucose and bone metabolism, body growth, and cardiovascular risk were analysed using a non-automated bridge assay protocol based on the method from EU patent 20170276675 together with Charité – Universitätsmedizin Berlin, Germany. Gliadin and tissue transglutaminase autoantibodies were measured by an automated Chemiluminescence Immunoassay (CLIA) from IDS (Immunodiagnostic Systems, Boldon, UK), see table S9. More autoantibodies are currently measured for endocrine and potentially cardiovascular receptors and targets of interest.

	Male (N=377)	Female (N=484)	Unit
Laboratory assessment	Mean ± SD	Mean ± SD	
Gliadin IgA	2.2 ± 4.9	0.7 ± 2.1	ng/ml
Gliadin IgG	2.0 ± 11.2	1.6 ± 11.4	ng/ml
Tissue transglutaminase IgA	2.5 ± 0.8	2.2 ± 0.5	ng/ml
Tissue transglutaminase IgG	0.3 ± 0.9	0.3 ± 1.1	ng/ml

Laboratory measurement: autoantibody phenotyping - abbreviations:

IgA = Immunoglobulin A, IgG = Immunoglobulin G

9. Metabolic phenotyping by Nuclear Magnetic Resonance (NMR)

Untargeted metabolomics was done in 1012 serum and urine samples using NMR-based metabolic phenotyping by NMR spectroscopy. Technical measurements were performed on a 600 MHz Avance Neo NMR spectrometer and using 1D CPMG (Carr–Purcell–Meiboom–Gill), NOESY and 2D J-resolved pulse sequences. Data processing: Principle Component Analysis (PCA) and Orthogonal Partial Least Squares Discriminant Analysis (O-PLS-DA) were performed. Metabolite reference chemical shifts from databases and metabolites were cross-checked using reference compounds and Chenomx software package if necessary. Metabolite concentrations were determined using internal/external standards and using the Eretic 2 approach implemented in Topspin [9,10].

10. Genome-wide characterization of cohort

More than 650,000 single nucleotide polymorphisms (SNPs) have been measured in a whole genome global screening array (GSA, Illumina bead chip (Infinium Global Screening Array-24 V2; Illumina Inc, USA)) in all cohort patients in cooperation with the Human Genotyping Facility (Genetic Lab at the Erasmus University Rotterdam, Netherlands) based on Illumina technologies in 2017. These genetic data, as well as specific genotyping e.g. for primary hypolactasia, are available for genome-wide association analyses and the inclusion in subsequent meta- and mega-analyses. As the results are stored in database, bioinformatical approaches such as diverse R-scripts allow for a fast gene-, pathway- or function-specific selection and export of SNP data. Epigenetic phenotyping is ongoing. The potential of various circulating non-coding RNAs for the diagnosis and prediction of (subclinical) diseases is currently evaluated for diabetes, as well as osteological, metabolic and CVD conditions.

11. Optometric phenotyping

For the ophthalmologic examinations, a Canon CR-2 AF (Canon Europa, Amstelveen, Netherlands) and for the optical coherence tomography (OCT) an OCT Spectralis (Heidelberg Engineering, Heidelberg, Germany) was used. Retinal vessel diameters were assessed with Integrative Vessel Analysis (IVAN software, N. Ferrier, University of Wisconsin, USA) and with the OCT device's inbuilt calliper tool from 12° peripapillary circle scans. Additional OCT scans included a macular volume scan and a high-resolution single scan through the fovea. Measured parameters are described in Table S10.

Table S10: Baseline parameters of optometric phenotyping		
	Mean ± SD	Unit
Right eye	Number (N=349)	
CRAE6	147.5 ± 12.5	µm
CRVE6	212.2 ± 17.7	µm
AVR	0.70 ± 0.06	
Left eye	Number (N=344)	
CRAE6	148.3 ± 13.6	µm
CRVE6	212.1 ± 18.1	µm
AVR	0.70 ± 0.06	

Optometric phenotyping – abbreviations:

CRAE6: mean diameter of the six biggest arterioles; CRVE6: mean diameter of the six biggest venules; AVR: arterioles-venules ratio

12. Biobank Storage

Until 2020, samples (Preservative: EDTA, Sodium-citrate and Lithium-heparin) were transferred immediately after sampling to an adjacent laboratory run by technicians from the local biobank. Sodium citrate samples were centrifuged at 2880g for 15min and the rest of the samples at 3360g for 10min at 4°C. The centrifuged biospecimen were then aliquoted manually. Since November 2013 samples were automatically aliquoted using a pipetting robot (Hamilton Microlab STARlet). After aliquoting, samples were temporally stored at -20°C. The transport to the final -80°C Biobank Graz storage unit was performed under ambient temperature. Since March 2019 the transport was performed under dry-ice conditions. Since 2020 these processes took place in another laboratory and technicians from the biobank picked up the samples at latest within one hour after sampling. Additionally, timepoints of sampling and execution of single work steps were digitally documented.

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