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NEW RESEARCH PAPER

Machine Learning-Derived Echocardiographic Phenotypes Predict Heart Failure Incidence in Asymptomatic Individuals

Masatake Kobayashi, MD, PhD,a,b,* Olivier Huttin, MD, PhD,a,b,* Martin Magnusson, MD, PhD,c,d,f João Pedro Ferreira, MD, PhD,a,b Erwan Bozec, PhD,a,b Anne-Cécile Huby, PhD,a,b* Gregoire Peud’homme, MSC,a,b Kevin Duarte, PhD,a,b Zohra Lamiral, MSC,a,b Kevin Dalleau, PhD,b Emmanuel Bresso, PhD,e Malika Smaïl-Tabbone, PhD,b,e Marie-Dominique Devignes, PhD,b,e Peter M. Nilsson, MD, PhD,c,e Margret Leosdottir, MD, PhD,c,d Jean-Marc Boivin, MD, PhD,a,b Faiez Zannad, MD, PhD,a,b Patrick Rossignol, MD, PhD,a,b Nicolas Girerd, MD, PhD,a,b on behalf of the STANISLAS study

ABSTRACT

OBJECTIVES This study sought to identify homogenous echocardiographic phenotypes in community-based cohorts and assess their association with outcomes.

BACKGROUND Asymptomatic cardiac dysfunction leads to a high risk of long-term cardiovascular morbidity and mortality; however, better echocardiographic classification of asymptomatic individuals remains a challenge.

METHODS Echocardiographic phenotypes were identified using K-means clustering in the first generation of the STANISLAS (Yearly non-invasive follow-up of Health status of Lorraine insured inhabitants) cohort (N = 827; mean age: 60 ± 5 years; men: 48%), and their associations with vascular function and circulating biomarkers were also assessed. These phenotypes were externally validated in the Malmö Preventive Project cohort (N = 1,394; mean age: 67 ± 6 years; men: 70%), and their associations with the composite of cardiovascular mortality (CVM) or heart failure hospitalization (HFH) were assessed as well.

RESULTS Three echocardiographic phenotypes were identified as "mostly normal (MN)" (n = 334), "diastolic changes (D)" (n = 323), and "diastolic changes with structural remodeling (D/S)" (n = 170). The D and D/S phenotypes had similar ages, body mass indices, cardiovascular risk factors, vascular impairments, and diastolic function changes. The D phenotype consisted mainly of women and featured increased levels of inflammatory biomarkers, whereas the D/S phenotype, consisted predominantly of men, displayed the highest values of left ventricular mass, volume, and remodeling biomarkers. The phenotypes were predicted based on a simple algorithm including $e^0$, left ventricular mass and volume ($e^0$VM algorithm). In the Malmö cohort, subgroups derived from $e^0$VM algorithm were significantly associated with a higher risk of CVM and HFH (adjusted HR in the D phenotype = 1.87; 95% CI: 1.04 to 3.37; adjusted HR in the D/S phenotype = 3.02; 95% CI: 1.71 to 5.34).

CONCLUSIONS Among asymptomatic, middle-aged individuals, echocardiographic data-driven classification based on the simple $e^0$VM algorithm identified profiles with different long-term HF risk. (4th Visit at 17 Years of Cohort STANISLAS-Stanislas Ancillary Study ESCIF [STANISLASV4]; NCT01391442) (J Am Coll Cardiol Img 2021;8:1010–1018) © 2021 The Authors. Published by Elsevier on behalf of the American College of Cardiology Foundation. This is an open access article under the CC BY-NC-ND license (http://creativecommons.org/licenses/by-nc-nd/4.0/).
A symptom left ventricular (LV) systolic dysfunction and ventricular enlargement or hypertrophy are classified under Stage B heart failure (HF), which is characterized by a high risk of developing HF or cardiovascular outcomes (1). Accumulating data have further broadened this category by incorporating LV diastolic dysfunction (DD) (2,3). These structural and functional cardiac alterations may vary with age and comorbidity burden such as hypertension and diabetes mellitus (4-8). Therefore, the inter-relationships of echocardiographic parameters can potentially complicate echocardiogram-based evaluations for identifying high-risk individuals in a population setting. Indeed, the definition of DD has been a moving target, and several classifications have been proposed over the last 20 years (9-12), leading to large differences in DD prevalence in population-based cohorts (13). Hence, a better use of echocardiographic phenotypes may help identify high-risk individuals and consequently prevent the progression from asymptomatic cardiac abnormalities to overt HF.

Machine learning (ML) clustering provides the potential value of identifying homogeneous groups among populations with heterogeneous underlying features. It has been efficiently used in various cardiovascular settings with noteworthy heterogeneity, eg, early cardiac structure alterations in diabetic patients (14) and patients with HF and preserved ejection fraction (HFpEF) (15). Given that ML clustering can identify groups of individuals with specific characteristics and risk, the relevance of those groups should consequently be apparent by their association with “hard” outcomes including HF onset and death, vascular profiles (associated with adverse cardiac structural and functional remodeling [16,17]) and circulating biomarkers associated with cardiovascular pathophysiology.

The authors hypothesized that echocardiographic phenotypes can be identified using ML-derived decision algorithms in asymptomatic populations and that those phenotypes are associated with different levels of vascular impairments, circulating biomarkers, and a long-term risk for HF. In keeping with the last impairment, the population-based STANISLAS (Yearly non-invasive follow-up of Health status of Lorraine insured inhabitants) cohort provided an opportunity to identify asymptomatic echocardiographic phenotypes and their associations with vascular parameters and proteomic profiles (18). Furthermore, the Malmö Preventive Project cohort provided the additional opportunity of verifying the external validity of this echocardiographic phenotyping by quantifying the association of these echocardiographic phenotypes with a long-term risk of incident HF and cardiovascular death (19).

METHODS

STANISLAS COHORT. The STANISLAS cohort was a single-center familial longitudinal population-based cohort from the Nancy region in France, the features of which have been previously described (18). The cohort, established in 1993 to 1995, enrolled families consisting of 2 parents and at least 2 biological children which were deemed healthy and initially free of declared illness, with the primary objective to investigate gene-gene and gene-environment interactions. Between 2011 and 2015, 1,705 participants underwent the 4th cohort visit. The study was approved by the local ethics committee (Comité de Protection des Personnes Est III-Nancy-France, 11.02.03), and all study participants gave informed consent to participate (NCT01391442).

Collected variables included age, sex, height, weight, smoking status, disease history, medication use, and the following cardiovascular risk factors: hypertension, diabetes mellitus, and dyslipidemia. Hypertension was diagnosed by 3 consecutive office blood pressure measurements or by the use of anti-hypertensive medication (20).

In the STANISLAS cohort, 701 participants who were initially children were relatively young at the 4th visit (mean age: 34 ± 5 years). As these participants are unlikely to have any echocardiographic abnormalities, only “parents” of the STANISLAS cohort were included in the present analysis. In
addition, participants with a history of HF or missing echocardiographic variables (used in the clustering procedure) were excluded (Supplemental Figure 1).

**CARDIAC FUNCTION ANALYSIS IN THE STANISLAS COHORT.** Echocardiographic examinations were performed at the Nancy University Hospital Clinical Investigation Center using standard, commercially available ultrasound equipment (Vivid 9, General Electric Medical Systems, Horten, Norway) with a 2.5-MHz phased-array transducer and reviewed on an Echopac workstation version 202 or 203 (GE Healthcare). The measurement of left atrial (LA) volume was obtained from the biplane method of disks (modified Simpson’s rule) using apical 4- and 2-chamber views at end-systolic phase (without dedicated LA views maximizing LA size).

Diastolic function was assessed from the mitral inflow pattern by pulsed-wave Doppler. Mitral annular early diastolic velocity (e’), was assessed at the septal and lateral sites of the mitral annulus using tissue Doppler imaging. The E/A ratio, e’ mean, and E/e’ mean ratio were calculated (10). LV systolic deformation was assessed from the 3 apical views with measurement of the global LV systolic strain (GLS %) corresponding to the maximum absolute value of strain derived from strain curves. DD was defined by the 2016 American Society of Echocardiography/European Association of Cardiovascular Imaging recommendations (DD class 2016 assessment with European Association of Cardiovascular Imaging the 2016 American Society of Echocardiography/Corresponding to the maximum absolute value of measurement of the global LV systolic strain (GLS %) information was assessed from the 3 apical views with Complior (Alam Medical) device (25). Intima media thickness (IMT) was measured using a linear probe with an echo-tracking system (Wall Track System [WTS]/ART.LAB) at 1 cm-2 cm below the carotid bifurcation on the right common carotid artery (26).

Circulating biomarkers expressing inflammation, extracellular matrix remodeling, and renal function. Inflammatory mediators associated with diastolic dysfunction in HFpEF were assessed (fatty acid-binding protein [FABP]-4 growth differentiation factor [GDF]-15, and galectin [Gal]-3). Adhesion molecules (vascular cell adhesion molecule [VCAM]-1 and intercellular adhesion molecule [ICAM]-1), transforming growth factor (TGF)-β-1, monocyte chemoattractant protein (MCP)-1, receptor for advanced glycosylation end (RAGE) products, interleukin (IL)-6, IL-8, IL-1 receptor-like 2 (IL1RL-2), tumor necrosis factor receptor 1 (TNF-R1), and osteopontin (OPN) were also studied as they were reported to be implicated in the inflammatory process related to HFpEF (28-33).

Extracellular matrix remodeling and angiogenesis in HFpEF were supported by the expression of matrix metalloproteinase (MMP)-1, MMP-2, MMP-9, MMP-12, N-terminal propeptide of procollagen type III (PIIINP), C-terminal propeptide of procollagen type I (PICP), soluble ST-2, N-terminal pro-B-type natriuretic peptide (NT-proBNP), brain natriuretic peptide (BNP), C-type natriuretic peptide (CNP), troponin-I, tissue inhibitor of metalloproteinase (TIMP)-1, TIMP-4, adrenomedullin (ADM), angiogenin (ANG), and vascular endothelial growth factor (VEGF)-A. Blood urea nitrogen (BUN) and creatinine were, furthermore, used to express renal function in HFpEF (34). These selected biomarkers were measured using Proseek (Olink Proteomics, Uppsala, Sweden) multiplex organ damage, metabolism, cardiovascular (CVD) II, III and Inflammation 96 × 96 kits in serum using proximity extension assay technology (35). In addition, PIIINP was measured by using radioimmunoassay (Orion Diagnostica, Eurofins Panlabs) and PICP by enzyme immunoassay (Metra, Quidel Corp.).

**MALMÖ PREVENTIVE PROJECT COHORT.** The Malmö Preventive Project cohort was a population-based longitudinal cohort of Malmö (Sweden) inhabitants (19). Of 1,792 participants having undergone an echocardiogram from 2002 to 2006, those without a
history of HF were included in our analysis (36). The Lund University’s ethics committee (Lund, Sweden) approved the study (LU 244-02), and written informed consent was obtained from each participant. HF hospitalization and other clinical outcomes were retrieved by record linkage between the Swedish personal identification number and national health care databases.

Echocardiograms were conducted using a 3V2c transducer (Acuson Sequoia, AGES-RS and MPP-RES) or an S3 transducer (Sonos 5500 Philips, MPP-RES). According to the recommendations, LV and LA structures, systolic and diastolic function were assessed (37,38). Intra- and interobserver variability and full description of the echocardiographic protocol have previously been reported (36) and are shown in Supplemental Table 1. The values of LV diastolic volume and LA volume were respectively estimated from LV diastolic diameter and LA area based on linear equations derived from the STANISLAS cohort (Supplemental Table 1). DD was coded according to the 2016 American Society of Echocardiography/European Association of Cardiovascular Imaging recommendations. All individuals were considered having a peak tricuspid regurgitation (TR) velocity <2.8 cm/s as this parameter was unavailable. In addition, individuals who had either DD, LVH, or an S3 transducer (Sonos 5500 Philips, MPP-RES) were retrieved by record linkage between the Swedish personal identification number and national health care databases.

MACHINE LEARNING APPROACH: CLUSTER ANALYSIS AND DECISION TREE CONSTRUCTION. The Proposed Requirements for Cardiovascular Imaging-Related Machine Learning Evaluation (PRIME) guideline (39) checklist is shown in Supplemental Table 2.

Cluster analysis was performed based on echocardiographic data (ie, LV mass index [LVMij], LV end-diastolic volume index [LVEDVi], LA volume index [LAVi], LVEF, GLS, E/A ratio, e’ mean, and E/e’ mean ratio).

K-means clustering analysis was performed to identify echocardiographic patterns using the K-means R program. Cluster analysis was performed on standardized values (mean = 0; SD = 1), based on the Euclidean distance as the dissimilarity metric. This algorithm aimed at finding clusters by: 1) initializing k points as the centroids; 2) assigning each data point to the closest centroid; and 3) recomputing the coordinates of the k centroids as the barycenters of the new clusters. This procedure was repeated until the centroids were stable. A total of 30 different quality measurements for different numbers of clusters were used to find the optimal number of clusters using the NbClust R program. Missing values were not imputed. In order to choose between available clustering methods, the same echocardiographic parameters were used in a hierarchical clustering and latent class model (LCM) clustering. K-means clustering achieved a higher internal consistency than the hierarchical clustering after cluster bootstrap (>1,000) but provided consistent results with the LCM approach (Supplemental Table 3). Thus, the authors reported the results of the K-means clustering as the primary analysis.

Decision trees provided parsimonious and transparent rules that were particularly appropriate in producing a hierarchy of 2 or more predictive factors simultaneously. Each patient in the data set (or the cohort) was labeled with a corresponding cluster. Decision trees were subsequently constructed to predict the clusters by using the rpart R program (40). This program selected the best threshold of each echocardiographic parameter to maximize the discrimination of clusters and evaluated through a 10-fold cross-validation to ensure that the prediction error was correctly estimated.

The overall accuracy of these cluster predictions was, furthermore, explored using random forest classification using the random Forest R application (41). Parameters of the random forest were optimized by a grid search with the Caret R package (42) by varying the number of trees between 50 and 1000, and modifying the number of selected variables to construct each split from 2 to the total number of available variables. Retained parameters were 500 trees and 15 random selected variables and the final model accuracy was evaluated by a 10-fold cross validation.

Principal component analysis was used to visualize the relationship between echocardiographic parameters used in cluster analysis and individuals in each echocardiographic phenotype.

The source code of all analyses performed within this project including the random forest model is available in GitLab.

STATISTICAL ANALYSIS. Categorical variables are described as frequencies (percentages), whereas continuous variables are described as mean ± SD or median interquartile ranges (IQR: 25th, 75th percentiles) depending on variable distributions. Comparisons of participant characteristics across echocardiographic phenotypes were analyzed using analysis of variance (ANOVA), Kruskal-Wallis, and chi-squared tests, as appropriate. Given the number of circulating biomarkers in the STANISLAS cohort, comparisons were analyzed using Kruskal-Wallis tests followed by p values corrected for the multiplicity of tests, using the Benjamini Hochberg method.
In the Malmö cohort, primary outcome was a composite of initial hospitalization for HF or cardiovascular death. Secondary outcome was a composite of cardiovascular mortality and hospitalization (HF, arrhythmia, or coronary artery disease). Survival probabilities were estimated using the Kaplan-Meier method. Cox proportional-hazards models for echocardiographic phenotypes were used to obtain unadjusted and covariate adjusted HRs. The covariates used for adjustments were Model 1: Heart OMics in Aging HF risk score (without creatinine) plus cystatin-C and NT-proBNP (43); Model 2: adjusted for ARIC (Atherosclerosis Risk In Communities) HF risk score without NT-proBNP; and Model 3: ARIC HF risk score plus NT-proBNP (44). The increased C index and net reclassification improvement (NRI) were implemented to assess added prognostic values of echocardiographic phenotypes for the primary outcome on top of: 1) ARIC HF risk score plus NT-proBNP; 2) cardiac structure and function abnormalities (ie, LVH, LAVi >7 cm²/m², septal e’ <9 cm/s or lateral e’ <10 cm/s, E/e’ mean >14); 3) DD class 2016 assessment with conventional echo variables; and 4) DD class 2016 assessment with additional echo biomarkers.

All statistical analyses were performed using R version 3.4.0 software (R Development Core Team). A value <0.05 was considered statistically significant.

RESULTS

Participants with missing values of at least 1 echocardiographic parameter used in cluster analysis were excluded (Supplemental Table 4). Those individuals

### Table 1: Clinical and Vascular Characteristics According to Echocardiographic Phenotypes in the STANISLAS Cohort

<table>
<thead>
<tr>
<th></th>
<th>Global (N = 827)</th>
<th>MN Phenotype (n = 334)</th>
<th>D Phenotype (n = 322)</th>
<th>D/S Phenotype (n = 170)</th>
<th>P Value</th>
<th>P Value for the Comparison Between D and D/S Phenotypes</th>
</tr>
</thead>
<tbody>
<tr>
<td>Age, yrs</td>
<td>60.0 ± 5.0</td>
<td>59.0 ± 4.0</td>
<td>61.0 ± 5.0</td>
<td>61.0 ± 5.0</td>
<td>&lt;0.001</td>
<td>0.11</td>
</tr>
<tr>
<td>Men</td>
<td>399.0 (48.2)</td>
<td>143.0 (42.8)</td>
<td>128.0 (39.6)</td>
<td>128.0 (75.3)</td>
<td>&lt;0.001</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>Body mass index, kg/m²</td>
<td>27.0 ± 4.0</td>
<td>25.0 ± 4.0</td>
<td>27.0 ± 4.0</td>
<td>28.0 ± 5.0</td>
<td>&lt;0.001</td>
<td>0.12</td>
</tr>
<tr>
<td>Medical history</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Hypertension</td>
<td>234.0 (28.3)</td>
<td>59.0 (17.7)</td>
<td>115.0 (35.6)</td>
<td>60.0 (35.3)</td>
<td>&lt;0.001</td>
<td>0.95</td>
</tr>
<tr>
<td>Diabetes</td>
<td>44.0 (5.4)</td>
<td>13.0 (3.9)</td>
<td>20.0 (6.3)</td>
<td>11.0 (6.5)</td>
<td>0.31</td>
<td>0.93</td>
</tr>
<tr>
<td>Dyslipidemia</td>
<td>259.0 (31.3)</td>
<td>81.0 (24.3)</td>
<td>118.0 (36.5)</td>
<td>60.0 (35.3)</td>
<td>0.001</td>
<td>0.79</td>
</tr>
<tr>
<td>Coronary artery disease</td>
<td>20.0 (2.4)</td>
<td>7.0 (2.1)</td>
<td>6.0 (1.9)</td>
<td>7.0 (4.1)</td>
<td>0.27</td>
<td>0.15</td>
</tr>
<tr>
<td>Smoking</td>
<td>101.0 (12.3)</td>
<td>42.0 (12.7)</td>
<td>45.0 (14.2)</td>
<td>14.0 (8.2)</td>
<td>0.16</td>
<td>0.06</td>
</tr>
<tr>
<td>Medications</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Antihypertensive treatment</td>
<td>225.0 (27.5)</td>
<td>58.0 (17.5)</td>
<td>104.0 (32.6)</td>
<td>63.0 (37.7)</td>
<td>&lt;0.001</td>
<td>0.26</td>
</tr>
<tr>
<td>Statin</td>
<td>156.0 (18.9)</td>
<td>40.0 (12.0)</td>
<td>76.0 (23.3)</td>
<td>40.0 (23.3)</td>
<td>&lt;0.001</td>
<td>0.99</td>
</tr>
<tr>
<td>ACC/AHA Heart Failure Stage</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Stage 0</td>
<td>398.0 (48.1)</td>
<td>216.0 (64.7)</td>
<td>151.0 (46.7)</td>
<td>31.0 (18.2)</td>
<td>&lt;0.001</td>
<td>0.81</td>
</tr>
<tr>
<td>Stage A</td>
<td>247.0 (25.9)</td>
<td>75.0 (22.5)</td>
<td>141.0 (43.7)</td>
<td>31.0 (18.2)</td>
<td>&lt;0.001</td>
<td></td>
</tr>
<tr>
<td>Stage B</td>
<td>182.0 (22.0)</td>
<td>43.0 (12.9)</td>
<td>31.0 (9.6)</td>
<td>108.0 (63.5)</td>
<td>&lt;0.001</td>
<td></td>
</tr>
<tr>
<td>Biochemistry</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>HbA1c, %</td>
<td>5.8 ± 0.6</td>
<td>5.7 ± 0.5</td>
<td>5.8 ± 0.6</td>
<td>5.8 ± 0.6</td>
<td>0.008</td>
<td>0.30</td>
</tr>
<tr>
<td>Total cholesterol, mg/dL</td>
<td>224.0 ± 38.0</td>
<td>225.0 ± 37.0</td>
<td>227.0 ± 39.0</td>
<td>218.0 ± 38.0</td>
<td>0.053</td>
<td>0.02</td>
</tr>
<tr>
<td>LDL cholesterol, mg/dL</td>
<td>142.0 ± 34.0</td>
<td>144.0 ± 33.0</td>
<td>143.0 ± 35.0</td>
<td>138.0 ± 34.0</td>
<td>0.25</td>
<td>0.31</td>
</tr>
<tr>
<td>eGFR, ml/min/1.73m²</td>
<td>91.0 ± 17.0</td>
<td>91.0 ± 16.0</td>
<td>91.0 ± 17.0</td>
<td>93.0 ± 17.0</td>
<td>0.48</td>
<td>0.18</td>
</tr>
<tr>
<td>Vascular parameters</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Heart rate, beats/min</td>
<td>62.0 ± 8.0</td>
<td>61.0 ± 8.0</td>
<td>63.0 ± 9.0</td>
<td>60.0 ± 8.0</td>
<td>&lt;0.001</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>Peripheral SP, mm Hg</td>
<td>124.0 ± 17.0</td>
<td>119.0 ± 15.0</td>
<td>126 ± 17</td>
<td>130 ± 18</td>
<td>0.044</td>
<td></td>
</tr>
<tr>
<td>Central SP, mm Hg</td>
<td>115.0 ± 17.0</td>
<td>110.0 ± 16.0</td>
<td>117.0 ± 17</td>
<td>119.0 ± 18</td>
<td>&lt;0.001</td>
<td>0.48</td>
</tr>
<tr>
<td>Peripheral AI, %</td>
<td>85.0 ± 17.0</td>
<td>85.0 ± 17.0</td>
<td>86.0 ± 17</td>
<td>82.0 ± 14</td>
<td>0.11</td>
<td>0.01</td>
</tr>
<tr>
<td>Central AI, %</td>
<td>34.0 ± 12.0</td>
<td>34.0 ± 11.0</td>
<td>35.0 ± 12</td>
<td>34.0 ± 13</td>
<td>0.30</td>
<td>0.29</td>
</tr>
<tr>
<td>IMT, μm</td>
<td>691.0 (616.0-784.0)</td>
<td>673.0 (598.0-764.0)</td>
<td>689.0 (627.0-771.0)</td>
<td>738.0 (656.0-822.0)</td>
<td>&lt;0.001</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>PWV, m/s</td>
<td>8.4 (7.5-9.4)</td>
<td>8.1 (7.3-9.0)</td>
<td>8.6 (7.7-9.6)</td>
<td>8.7 (7.6-9)</td>
<td>0.001</td>
<td>0.56</td>
</tr>
<tr>
<td>TAC index, ml/mm Hg/m²</td>
<td>1.00 (0.79-1.25)</td>
<td>1.06 (0.88-1.32)</td>
<td>0.95 (0.73-1.17)</td>
<td>1.01 (0.77-1.33)</td>
<td>&lt;0.001</td>
<td>0.03</td>
</tr>
<tr>
<td>SVR index, dyne x s/cm²/m²</td>
<td>2,624.0 (2,254.0-3,148.0)</td>
<td>2,568.0 (2,237.0-3,007.0)</td>
<td>2,648.0 (2,299.0-3,235.0)</td>
<td>2,724.0 (2,223.0-3,360.0)</td>
<td>0.021</td>
<td>0.83</td>
</tr>
</tbody>
</table>

Values are mean ± SD, n (%), or median interquartile range (25th, 75th percentiles). Bold values indicate statistical significance.

**ACC/AHA** = American College of Cardiology/American Heart Association; AI = augmentation index; D = diastolic changes; D/S = diastolic changes with structural remodeling; eGFR = estimated glomerular filtration rate; HbA1c = glycylated hemoglobin; IMT = intima media thickness; IQR = interquartile range; LDL = low-density lipoprotein; PWV = pulse wave velocity; SP = systolic pressure; STANISLAS = Yearly non-invasive follow-up of Health status of Lorraine insured inhabitants; SVR = systemic vascular resistance; TAC = total arterial compliance.

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more often had comorbidities (eg, obesity, hypertension, diabetes) than those with complete echocardiographic data. In a total of 827 participants included in the current analysis 60 ± 5 years (48.2% men), mean body mass index was 27 kg/m², 28.3% of participants had hypertension, 5.4% had diabetes mellitus, and mean LVEF was 66% (Tables 1 and 2).

In the Malmö Preventive Project Cohort, a total of 1,394 participants were included (Table 3). Age was 67 ± 6 years, 70.0% of participants were men, mean body mass index was 28 kg/m², 63.7% had hypertension, and 34.3% had diabetes mellitus.

**Clusters Derived from Echocardiographic Data.** The present cluster analysis identified 3
groups with different echocardiographic phenotypes (Tables 1 and 2, Figure 1, Supplemental Figure 2). The largest cluster (n = 334), labeled mostly normal (MN) phenotype, had the highest e′ and E/A ratio and the highest (absolute) LV systolic strain. The next largest cluster (n = 323), labeled “diastolic changes (D)” phenotype, had lower e′ and higher E/e′ ratio. The last cluster (n = 170), labeled “diastolic changes with structural remodeling (D/S)” phenotype, had the highest LV mass and volumes, highest LA volume, and lowest (absolute) LV systolic strain as well as lower e′ and higher E/e′ ratio. The prevalence of DD class 2016 assessment with conventional echo biomarkers was very low in all 3 phenotypes (<5%), and approximately one-half of individuals with D/S phenotype had DD class 2016 assessment with additional echo biomarkers (Table 2).

Participant characteristics according to hierarchical clustering-derived echocardiographic phenotypes are presented, and similar patterns of echocardiographic phenotypes were observed between K-means clustering and hierarchical clustering (Supplemental Table 5, Supplemental Figure 3). In addition, the LCM model provided the consistent phenotypes with the K-means clustering (Rand index: 0.83) (Supplemental Table 3).
When performing cluster analysis separately in men and women, similar patterns of echocardiographic phenotypes (ie, phenotypes D and D/S) were observed (Supplemental Figure 3).

**CLINICAL CHARACTERISTICS ACCORDING TO ECHOCARDIOGRAPHIC PHENOTYPES IN THE STANISLAS COHORT.** The D phenotype consisted mainly of women, whereas the D/S phenotype included predominantly of men. The D and D/S phenotypes exhibited older age, higher body mass index, and more often cardiovascular risk factors than the MN phenotype; however, no significant differences among these clinical features were observed between D and D/S phenotypes (Table 1).

**VASCULAR PROFILES ACCORDING TO ECHOCARDIOGRAPHIC PHENOTYPES IN THE STANISLAS COHORT.** The MN phenotype had more favorable levels of central and peripheral pressure and vascular stiffness (PWV and IMT) than the D and D/S phenotypes, whereas no significant differences were observed between the D and D/S phenotypes (PWV [m/s], 8.6 [95% CI: 7.7 to 9.6] in D phenotype vs 8.7 [95% CI: 7.6 to 9.8] in D/S phenotype; \( P = 0.56 \)) (Table 1, Figure 1).

**CIRCULATING BIOMARKERS ACCORDING TO ECHOCARDIOGRAPHIC PHENOTYPES IN THE STANISLAS COHORT.** Of the 32 circulating biomarkers, 14 were significantly different across the 3 phenotypes. These
biomarkers were associated with different pathophysiological domains (ie, inflammation and extracellular matrix remodeling) (Figure 2, Supplemental Table 6). Overall, the D phenotype exhibited the highest levels of circulating biomarkers associated with inflammation (ie, FABP4, IL-6, and IL1RL2). Markers generally associated with remodeling (ie, GDF15, PIIINP, ST-2, troponin-I, and CNP, increased from the MN to the D and D/S phenotypes, with the highest ST-2, troponin-I, and CNP values observed in the D/S phenotype.

DECISION TREE AND RANDOM FOREST TO IDENTIFY ECHOCARDIOGRAPHIC PHENOTYPES IN THE STANISLAS COHORT: “e’VM” ALGORITHM. The decision tree identified mean ’e’, LVEDVi, and LVMi as the most relevant variables to correctly classify participants in 1 of the echocardiographic profiles. The global accuracy of the decision tree was good (79.0%) (Figure 3). Addition of clinical variables (ie, age, sex, body mass index, prevalence of hypertension, diabetes, dyslipidemia, coronary artery disease, and smoking) to the candidate variables for decision tree building did not modify the decision algorithm; hence, these clinical variables were not retained by the algorithm as relevant additions to the decision tree. In the following sections, the authors refer to the constructed decision tree as the “e’VM” algorithm. Furthermore, the random forest classifier (available at Gitlab software) for these echocardiographic phenogroups provided an overall accuracy of 0.86.

EXTERNAL VALIDATION OF e’VM ALGORITHM IN THE MALMÖ PREVENTIVE PROJECT COHORT. Participant characteristics according to e’VM algorithm phenotypes in the Malmö cohort are presented in Table 3. The MN phenotype (n = 440) displayed the most favorable values of clinical, biological, and echocardiographic profiles. The D phenotype
included 512 participants and displayed the lowest proportion of men, whereas the D/S phenotype (n = 442) featured a high proportion of men, the highest NT-proBNP level, and the largest LV mass and LA surface area indices. Similar age, body mass index, cardiovascular risk factors and diseases (ie, hypertension, dyslipidemia and renal function) and diastolic function alterations were observed in the D and D/S phenotypes (Table 3).

**SURVIVAL ANALYSIS IN THE Malmö Preventive Project Cohort.** In the Malmö study, during a median follow-up of 10.3 years (9.8 to 11.1 years), 10.1% of participants met the primary outcome. Compared with the MN phenotype, both the D and D/S phenotypes were significantly associated with increased rates of the primary outcome (crude HR: 2.47; 95% CI: 1.38 to 4.41; P = 0.002 for the D phenotype and crude HR: 4.67; 95% CI: 2.67 to 8.14; P < 0.001 for the D/S phenotype) (Figure 4, Supplemental Table 7). After adjustment for the ARIC HF risk score plus NT-proBNP, the D and D/S phenotypes remained significantly associated with the primary outcome (adjusted HR: 1.87; 95% CI: 1.04 to 3.37; P = 0.03 for the D phenotype and adjusted HR: 3.02; 95% CI: 1.71 to 5.34; P < 0.001 for the D/S phenotype). Poor prognosis of the D/S phenotype persisted even in terms of the composite of cardiovascular hospitalization and mortality (Supplemental Table 7).

**DISCUSSION**

The present analysis enabled identifying 3 echocardiographic phenotypes with differential biological characteristics and risk of HF, namely, “mostly normal (MN),” “diastolic changes,” and “diastolic changes with structural remodeling” phenotypes (Central Illustration). The D phenotype had the most abnormal diastolic function parameters and typically exhibited higher levels of inflammatory biomarkers, whereas the D/S phenotype expressed LV and LA remodeling as well as diastolic changes and was associated with higher levels of extracellular matrix remodeling. Both phenotypes were associated with vascular impairments as well as a higher risk of HF hospitalization and/or cardiovascular mortality. Our findings suggest that the easy-to-use e’VM echocardiographic-based algorithm presented herein may identify asymptomatic individuals with a higher
Importantly, this eVM algorithm was validated externally in an independent Swedish cohort, which suggests that this algorithm could be widely applicable in White populations.

**ECHOCARDIOGRAPHIC PHENOTYPES.** In the current study, LVEF and DD prevalence of the participants were similar to those of the general populations described in previous reports (45–48). Of note, in the present classification, both LVEDVI and LVM were key components in classifying asymptomatic individuals into the proposed echocardiographic phenotypes. By contrast, GLS was not selected as a key marker for the eVM algorithm despite its well-known performance to detect early alterations of LV systolic function in a range of populations (49,50). The echocardiographic phenotypes identified in the present analysis were notably similar to the phenotypic patterns of early cardiac alterations reported in diabetic patients (14). Indeed, using a cluster analysis based on echocardiographic parameters in 842 diabetic patients, Ernande et al. (14) highlighted 3 echocardiographic phenotypes including a “diastolic dysfunction” phenotype and “LVH with systolic dysfunction” phenotype. Despite the differing settings (general population vs diabetic patients), these findings would appear to reinforce our results and the possible ubiquitous nature of the echocardiographic phenotypes based on early and/or asymptomatic cardiac dysfunction. Furthermore, the prevalence of DD class 2016 assessment with conventional echo variables was very low in our cohort and remained similar in both D and D/S phenotypes, suggesting that the current definition of DD may be less relevant in an asymptomatic population.

The D phenotype was more frequent in women, whereas men were more likely to express diastolic changes with structural remodeling. The results reported by Ernande et al. were perceived as sex-specific echocardiographic patterns (14,51). However, our analysis showed similar patterns of echocardiographic phenotypes derived from men and women subgroups. The echocardiographic phenotypes proposed herein could thus provide robust identification of individuals regardless of sex difference.

Arterial stiffness is a fundamental contributor to increasing diastolic pressure (16,17) as well as impairing early systolic function (52). In the present study, (proximal to distal) vascular deteriorations were observed in both the D and the D/S phenotypes. Of note, PWV was similar in these 2 phenotypes. Underlying mechanisms following vascular abnormalities may potentially occur in these 2 phenotypes: the D phenotype may exhibit an early increase in intracardiac pressures in the absence of structural remodeling, while the D/S phenotype may further compensate the increase in afterload using increased LV volume and mass. This hypothesis could shed new
light on the early stage of the inter-relationship between cardiac and vascular parameters.

**ECHOCARDIOGRAPHIC PHENOTYPES AND CIRCULATING BIOMARKERS ASSOCIATED WITH INFLAMMATION OR EXTRACELLULAR MATRIX REMODELING.** Inflammation and extracellular matrix remodeling have been found to be associated with diastolic function (30,53). Pau-lus and Tschope (29) proposed a paradigm for HFpEF which highlighted the importance of a systemic proinflammatory state related to ageing, hypertension and obesity, further leading to the development of DD. In the current analysis, our results show that the D phenotype was significantly associated with increased levels of inflammatory biomarkers such as FABP4. Indeed, FABP4 is mainly expressed in adipose tissue and is an important mediator of inflammation, showing a significant link with DD (54,55) and a higher risk of cardiovascular events (56).

The D/S phenotype, on the other hand, was significantly associated with increased levels of extracellular matrix remodeling biomarkers. Both echocardiographic phenotypes were associated with a higher risk of developing heart failure or cardiovascular death. CV = cardiovascular; HF = heart failure; IL1RL2 = Interleukin 1 Receptor Like 2; LVDVi = LVDV/ left ventricle diastolic volume index; LVMi = left ventricular mass index; ST2 = soluble interleukin 2.


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**CENTRAL ILLUSTRATION** Echocardiographic Phenotypes in Asymptomatic Individuals

<table>
<thead>
<tr>
<th>Mostly Normal</th>
<th>Diastolic Changes</th>
<th>Diastolic Changes with Structural Remodeling</th>
</tr>
</thead>
<tbody>
<tr>
<td>e’ ≥10</td>
<td>e’ &lt;10</td>
<td>e’ &lt;10</td>
</tr>
<tr>
<td>LVDVi &lt;50 + LVMi &lt;105</td>
<td>OR</td>
<td>LVDVi ≥50 + LVMi &lt;85</td>
</tr>
<tr>
<td>LVDVi ≥50 + LVMi ≥105</td>
<td>OR</td>
<td>LVDVi ≥50 + LVMi ≥85</td>
</tr>
</tbody>
</table>

**Predominantly women**

- Inflammation markers:
  - Fatty acid-binding protein-4
  - Interleukin-6
  - IL1RL2

**× 2 risk**

HF and/or CV death compared with mostly normal

**Predominantly men**

- Remodeling markers:
  - Soluble ST-2
  - Troponin-I
  - C-type natriuretic peptide

**× 5 risk**

HF and/or CV death compared with mostly normal

Among the 3 echocardiographic phenotypes, the e’VM echocardiographic-based algorithm (ie, mean e’, left ventricular volume, and mass) differentially identified asymptomatic individuals with the “diastolic changes” phenotype and those with the “diastolic changes with structural remodeling” phenotype. The “diastolic changes” phenotype, consisting predominantly of women, displayed higher circulating levels of inflammatory biomarkers, whereas the “diastolic changes with structural remodeling” phenotype, consisted predominantly of men, displayed higher circulating levels of extracellular matrix remodeling biomarkers. Both echocardiographic phenotypes were associated with a higher risk of developing heart failure or cardiovascular death. CV = cardiovascular; HF = heart failure; IL1RL2 = Interleukin 1 Receptor Like 2; LVDVi = LVDV/ left ventricle diastolic volume index; LVMi = left ventricular mass index; ST2 = soluble interleukin 2.
Our results, hence, support the fact that systemic inflammation may be associated with rising LV pressure with changes in diastolic function, whereas microvascular endothelial dysfunction and myocardial damage may be more closely related to cardiac structural remodeling, even in asymptomatic individuals.

**CLINICAL IMPLICATIONS.** In the latest American College of Cardiology/American Heart Association guidelines, an echocardiogram screening is recommended to identify those at high-risk of developing HF in individuals with hypertension (59) or slightly increased natriuretic peptide level (BNP: >50 pg/mL [Class IIa to IIb recommendation]) (60). Importantly, we show that the simple eVM echocardiographic algorithm improved risk stratification on top of currently available scores such as the ARIC HF risk score and traditionally defined cardiac structure/function abnormalities (including DD class 2016 assessment), thus suggesting that additional echocardiogram-based screening could be encouraged. Of further importance, among the presented “high-risk” subgroups, DD class 2016 assessment with conventional echo variables only diagnosed <5% of individuals. Our results consequently suggest the importance of re-assessing echocardiographic data using the eVM algorithm despite the absence of cardiac dysfunction according to conventional recommendations. We believe our echocardiographic phenotypes may serve as a new evidence-driven, integrated classification of individuals in preclinical settings with plausible circulating biomarkers and notable prognostic implications. How to manage individuals identified at high risk based on our algorithms is yet unknown. At least, given the higher risk of these individuals, a more careful follow-up could be undertaken. Whether the use of cardiovascular drugs could reduce the occurrence of HF in asymptomatic individuals requires testing. Importantly, the random forest model we built could be imbedded in health record systems to provide an automated identification of high-risk patients, maximizing the usefulness of echocardiogram in asymptomatic individuals.

**STUDY LIMITATIONS.** The main limitation of this study is its inherent observational design, based on a cross-sectional assessment; causality cannot therefore be inferred. The number of individuals with incident HF was relatively low, leading to moderate statistical power. The eVM algorithm was derived from only echocardiographic variables; however, when adding additional clinical variables (ie, age, sex, comorbidities), the algorithm remained unchanged, suggesting that echocardiographic variables alone better predicted our phenotypes. In addition, the decision tree had numerically lower accuracy than the random forest, but this difference was moderate. This may be caused by some overoptimism in the decision tree accuracy statistics, as random forest are expected to perform sizably better than decision trees. As the cohort included relatively young individuals with a low comorbidity burden, the generalization of this phenotypes should be limited to individuals aged 50 to 70 years without overt cardiovascular disease. Further research is warranted to explore the applicability of our echocardiogram-based algorithm in older and sicker populations. In the STANISLAS cohort, individuals who had missing values for echocardiographic parameters used in cluster analysis were excluded. These latter individuals more often had cardiovascular risk factors. Echocardiographic data were acquired at different time periods in the STANISLAS and Malmö cohorts. However, the recommended measurement of echocardiographic parameters has not changed in the last 20 years (9–12). In the Malmö cohort, peak TR velocity was not quantified and was considered >2.8 cm/s as only a few individuals (<2%) have been reported to have a peak TR velocity greater than >2.8 cm/s in a similar population-based cohort (45). Importantly, in the Malmö cohort, LV end-diastolic volume and LA volume were not quantified; hence, these values were respectively estimated from LV diastolic diameter and LA surface area using a linear equation. This important limitation should trigger other external validations in prospective multicenter studies to improve the generalization of the eVM algorithm. However, despite this limitation, the eVM algorithm predicted outcomes in the Malmo cohort.

**CONCLUSIONS**

The present analysis allowed highlighting 3 echocardiographic phenotypes than can be easily identified in clinical practice using a simple algorithm (the eVM algorithm). The 2 most abnormal phenotypes, namely “diastolic changes” and “diastolic changes with structural remodeling,” exhibited significant differences in circulating biomarkers and the risk of HF and/or cardiovascular death. The authors believe that these echocardiographic phenotypes shed new light on our understanding of asymptomatic cardiac dysfunction, and our findings may have significant clinical implications in the design of HF prevention strategies. A further prospective multicenter study is
needed to assess the applicability of this eVM algorithm.


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ADDRESS FOR CORRESPONDENCE: Prof Nicolas Girerd, Centre d’Investigation Clinique Pierre Drouin-INSERM-CHRU de Nancy, Institut lorrain du coeur et des vaisseaux Louis Mathieu, 4, rue du Morvan, 54500 Vandoeuvre-Les-Nancy, France. E-mail: n.girerd@chru-nancy.fr.
In asymptomatic individuals, a simple echocardiographic data-driven algorithm, based on e’, and LV volume and mass allows for identifying echocardiographic phenotypes with a high risk of developing HF.


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**KEY WORDS** biomarkers, cardiovascular diseases, cluster analysis, echocardiogram, heart failure, machine learning, prognosis

**APPENDIX** For supplemental figures and tables, please see the online version of this paper.