

Risk prediction in patients with heart failure with preserved ejection fraction: the LIFE-Preserved model

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13 ABSTRACT

14 **Background and aims** Heart failure (HF) with preserved ejection fraction (HFpEF) constitutes a
15 heterogeneous disease with varying prognosis. Given the rising incidence of HFpEF, accurate risk
16 prediction for these patients is needed to identify high-risk individuals, who may benefit the most
17 from preventive treatments. The LIFE-Preserved model was developed and validated for the
18 prediction of individual short-term and lifetime risk for HF hospitalization or cardiovascular (CV)
19 death in patients with HFpEF.

20 **Methods** LIFE-Preserved was derived in 20,332 patients aged 40-90 years with a left ventricular
21 ejection fraction $\geq 50\%$ from the Swedish HF Registry (SwedeHF). Cause- and sex-specific Cox
22 models were derived to predict the risk of HF hospitalization or CV death using 14 routinely
23 available predictors. Use of age as the timescale allowed for predictions beyond the maximum
24 follow-up duration in the derivation data, adjusted for competing risks. External validation was
25 performed in two trials (EMPEROR-Preserved, TOPCAT-Americas) and three registries (NHS
26 England Secure Data Environment, Veterans Affairs, HF-Particles). Model performance was
27 assessed by discrimination and calibration.

1 **Results** During a median follow-up of 1.8 years (interquartile range 0.6 – 4.2, maximum 19 years),
2 9341 first HF hospitalizations or CV deaths (46%) were observed in SwedeHF. External validation
3 included data from 28 062 patients with HFpEF (9930 [35%] first HF hospitalizations or CV deaths).
4 Pooled C-statistics were 0.714 (95% confidence interval [CI] 0.652–0.775) in trials and 0.658 (95%
5 CI 0.599–0.717) in registries, with adequate calibration in all external validation sources.
6 Performance was similar in men and women. An interactive calculator of the LIFE-Preserved model
7 has been made available here.

8 **Conclusions** The LIFE-Preserved model enables prediction of short-term and lifetime risk of HF
9 hospitalization or CV death in patients with HFpEF. The model could serve as a tool to identify high-
10 risk HFpEF patients, guiding clinical management and shared decision-making.

11

12 INTRODUCTION

13 The incidence and prevalence of heart failure (HF) with preserved ejection fraction (HFpEF)
14 is rising globally due to the ageing population and increasing cardiometabolic risk factors.^{1,2} HFpEF
15 is associated with high mortality rates, frequent hospitalizations and significant healthcare costs.¹⁻
16 ³ In recent years, new therapies have emerged that improve outcomes in patients with HFpEF.⁴⁻⁷
17 Sodium–glucose cotransporter 2 (SGLT2) inhibitors and the non-steroidal mineralocorticoid
18 receptor antagonist finerenone reduce the composite risk of HF hospitalizations and
19 cardiovascular (CV) death in patients with HFpEF.⁷⁻⁹ Semaglutide, a glucagon-like peptide-1 (GLP-
20 1) receptor agonist, and tirzepatide, a dual GLP-1 and glucose-dependent insulintropic
21 polypeptide (GIP) agonist, reduce the risk of HF hospitalization and CV death in patients with
22 obesity-related HFpEF.^{10,11} However, patients with HFpEF have highly heterogeneous clinical

1 profiles with varying prognosis, and absolute benefit of treatment options may vary greatly between
2 patients.¹² In fact, some patients are at relatively low risk, resulting in reduced absolute risk
3 reductions, a higher number needed to treat (NNT), and potentially lower cost-effectiveness.⁸ This
4 highlights the need for accurate risk prediction to optimize treatment allocation to patients most
5 likely to benefit. In addition, risk prediction may facilitate shared decision-making and increase
6 motivation for drug treatment and lifestyle changes.¹³

7 Previous prediction models for the HFpEF population are restricted to predict 1- or 2-year
8 risks.¹⁴⁻¹⁹ However, as HFpEF is a chronic disease requiring lifelong treatment, prediction models
9 should ideally provide long-term or lifetime risk to align with the clinical decision-making horizon.²⁰
10 Additionally, prior models were not HFpEF specific or were derived only in trial populations, which
11 are known to suffer from healthy participant bias. Therefore, to ensure clinical applicability,
12 validation in real-world population-based data is required. The European Society of Cardiology
13 (ESC) CV disease prevention guidelines recommend the use of 10-year and lifetime risk predictions
14 to guide clinical decision-making.²¹⁻²⁴ For patients with HF and reduced ejection fraction (HFrEF),
15 the LIFE-HF model is an externally validated risk prediction model for the estimation of lifetime risk
16 of HF hospitalization or CV death.²⁵ A similar model for patients with HFpEF could help to improve
17 HFpEF management by supporting personalized medicine and shared decision-making.

18 The objective of the present study was to develop and externally validate the LIFE-Preserved
19 model for individual prediction of short-term and lifetime risk of HF hospitalization or CV death in
20 patients with HFpEF.

21

1 METHODS

2 Study populations

3 The LIFE-Preserved risk model was developed in HFpEF patients from the Swedish HF
4 Registry (SwedeHF). Details on SwedeHF have been published previously.²⁶ Briefly, SwedeHF is an
5 ongoing nationwide registry in Sweden founded in 2000, with nationwide coverage since 2003.
6 Around 85% of all hospitals in Sweden actively participate in the registry and in 2020 approximately
7 31% of all prevalent HF patients in Sweden were registered.^{27,28} SwedeHF includes patients based
8 on a clinician-judged diagnosis of HF, and, since 2017, also identifies patients through the National
9 Patient Register using International Classification of Diseases 10th revision (ICD-10) codes (I50.0,
10 I50.1, I50.9, I42.0, I42.6, I42.7, I25.5, I11.0, I13.0, I13.2). Patients can be included from the
11 outpatient and the inpatient setting in SwedeHF. Baseline characteristics are recorded upon
12 inclusion. Information regarding medical history and other variables (e.g. comorbidities, date and
13 cause of death, hospitalizations, socioeconomic status, and prescriptions) can be retrieved
14 through linkage with the Swedish National Patient Register, Cause of Death Register, Statistics
15 Sweden and National Prescribed Drug Register (Supplementary Table S1). Linkage is achieved
16 through the personal unique identification number which all the residents in Sweden have.

17 For the current study patients aged between 40 and 90 years, with a left ventricular ejection
18 fraction (LVEF) $\geq 50\%$ at baseline and no earlier registration with documented LVEF $< 50\%$ were
19 eligible for analyses. Only those patients with a registration in SwedeHF between January 1st 2006
20 and February 1st 2020 were included to eliminate the potential impact of the COVID-19 pandemic
21 on study results and follow-up was truncated on February 1st 2020 for the same reason. If a patient
22 had multiple eligible registrations, the first one was selected (see Supplementary Figure S1 for the
23 flowchart of patient selection).

1 External validation was performed in HFpEF (LVEF \geq 50%) patients using population-based data
2 from three registries, the HF-Particles cohort in Poland²⁹, the Department of Veterans Affairs cohort
3 in the United States of America (VA)³⁰ and the NHS England Secure Data Environment (NHSE SDE)
4 through the British Heart Foundation Data Science Centre CVD-COVID-UK/COVID-IMPACT
5 Consortium³. In addition, external validation was performed in HFpEF (LVEF \geq 50%) patients using
6 data from two trials, the EMPEROR-Preserved trial⁷ and TOPCAT-Americas trial.³¹ Patients aged 40–
7 89 years were included in the external validation analyses, as the LIFE-Preserved model estimates
8 risks up to age 90, allowing 1-year risk prediction for individuals aged 89 years. Detailed
9 descriptions of the external validation data sources can be found in the Supplementary Methods.

11 Outcomes and predictors

12 The primary outcome was a composite of HF hospitalization or CV death, which was
13 assessed with a time-to-first-event analysis. This endpoint was chosen because it represents a
14 clinically actionable outcome in HFpEF, directly targeted by contemporary preventive therapies,
15 and therefore most relevant for informing risk stratification and clinical decision making. Non-CV
16 death was used as the competing outcome.

17 In SwedeHF, hospitalizations were extracted from the National Patient Register and HF
18 hospitalization was defined as the first hospitalization with HF registered as primary diagnosis
19 during follow-up. CV death was defined as death with CV disease registered as the primary cause
20 of death in the Cause of Death Register. All deaths that were not classified as CV deaths were
21 defined as non-CV deaths. In external validation, outcomes were assessed by linkage to hospital
22 records and mortality registries (the NHSE SDE, HF-Particles and VA) or adjudicated by an endpoint
23 committee in clinical trials (EMPEROR-Preserved and TOPCAT-Americas). Full details on outcome

1 definitions and the corresponding ICD-10 codes for all data sources are described in
2 Supplementary Table S2.

3 Candidate predictors of the LIFE-Preserved model were based on a literature review of prior
4 risk models (Supplementary methods and Figure S2).¹⁴⁻¹⁹ Candidate predictors were: diabetes
5 mellitus, current smoking, any previous hospitalization for HF, chronic obstructive pulmonary
6 disease (COPD), New York Heart Association (NYHA) class, ischaemic heart disease, N-terminal
7 pro-B-type natriuretic peptide (NT-proBNP), body mass index (BMI), heart rate, haemoglobin, and
8 estimated glomerular filtration rate (eGFR) based on serum creatinine calculated with the 2009
9 Chronic Kidney Disease Epidemiology Collaboratopn formula³² (Supplementary Table S3).
10 Additionally, an expert panel reviewed predictors based on clinical relevance and availability for
11 HFpEF and five additional predictors were identified for consideration: atrial fibrillation,
12 cerebrovascular disease (stroke or transient ischaemic attack), loop diuretic use, systolic blood
13 pressure, and duration of HF. Of these, only history of atrial fibrillation showed a statistically
14 significant improvement of internal C-statistics for the primary outcome of HF hospitalization or CV
15 death compared to the core model, and was therefore included in the final model (Supplementary
16 Figure S3). The same predictors were used for the primary and the competing outcomes.

17 Missing data were handled with imputation, using predictive mean matching for continuous
18 variables and logistic regression for binary variables. For predictor variables with missing data in
19 SwedeHF, values were imputed using two separate sex-specific multiple imputation by chained
20 equations models with 45 imputed datasets and 45 iterations, approximately the maximum
21 predictor missingness (Supplementary Table S4 and details in Supplementary Methods).
22 Convergence of the imputation models was checked by inspecting trace plots. Missing predictor
23 values in the external validation data were handled within each data source separately
24 (Supplementary Methods).

1

2 Model development

3 To allow for sex-differences in the relative effects of predictors, cause-specific Cox
4 proportional hazards models were derived for men and women separately. For each sex, two
5 separate cause-specific Cox proportional hazards models were constructed for prediction of first
6 hospitalization for HF or CV death and prediction of non-CV death as a competing outcome.
7 Adjusting for competing events is necessary for accurate risk prediction in a high-risk population to
8 avoid overestimation of risk estimates.³³ Age was used as the time scale with age at inclusion
9 defined as the start of follow-up (left truncation) and age at event or censoring defined as the end of
10 follow-up (right censoring). This allows for estimation of risks over the age range of the entire
11 population, enabling estimation risk at any time horizon as well as lifetime risk (i.e. risk until age
12 90).³⁴⁻³⁶ To account for the high readmission risk observed in recently hospitalized HFpEF patients,
13 a time-varying first-year risk adjustment was applied to patients with a recent HF hospitalization
14 (within 6 months prior to inclusion).³⁷ Continuous predictors were truncated at the 1st and 99th
15 percentiles to limit outlier effects, and non-linear transformations were applied if they improved
16 model fit based on Akaike's Information Criterion (AIC), details in Supplementary Methods. Age
17 interactions were considered when the proportional hazards assumption was violated, as
18 assessed by visual inspection of plotted Schoenfeld residuals. As SGLT2 inhibitors and GLP-1
19 receptor agonists were hardly used during the study period (<1% use at baseline), these were not
20 accounted for in model development.

21 Sex-specific baseline survival probabilities were derived for HF hospitalization or CV death
22 and non-CV death by estimating 3-month interval survival over the complete age range (40-90
23 years). All baseline survival probabilities were estimated for a reference patient and smoothed

1 using non-linear functions weighted by the number of patients in each age interval (Supplementary
2 Methods).

3

4 Individual and lifetime risk prediction

5 For each patient, the risk of HF hospitalization or CV death and non-CV death was
6 estimated for every remaining 3-month interval up to age 90 using validated lifetable methods.³⁴⁻³⁶
7 This approach, which has been shown reliable for prediction horizons of at least 17 years,³⁴
8 provides interval-specific survival probabilities for both outcomes. These were combined to
9 calculate the probability of surviving each interval free of events, accounting for competing risks.
10 These interval survival probabilities were multiplied to calculate the cumulative probability of
11 reaching a specific age free of any event. HF hospitalization-free life expectancy was defined as the
12 age at which this cumulative survival probability drops below 50%. Lifetime risks for each outcome
13 were defined as the cumulative risk from the current age of the patient until the maximum age of 90
14 years. Detailed descriptions of lifetable estimations are provided in the Supplementary Methods.

15

16 External validation

17 Performance of the LIFE-Preserved model was assessed by estimating Harrell's C-statistic
18 for discrimination and calibration plots of the predicted versus observed risk for calibration. All
19 were adjusted for competing risks.³⁸ To ensure stable estimates, external validation was performed
20 on the last full year in which at least 75th percent was still in follow-up in the respective data
21 source, which was at 2 years for EMPEROR-Preserved (n = 3928) and NHSE SDE (n = 28 375), 4
22 years for TOPCAT-Americas (n = 1495), and 8 years for HF-Particles (n = 838). In the VA, which was

1 the largest cohort with long term follow-up, external validation was performed at 10 years (n =
2 4499). C-statistics were pooled using inverse-variance weighting³⁹, which was performed
3 separately for trials (EMPEROR-Preserved and TOPCAT-Americas) and registries (NHSE SDE, HF-
4 Particles and VA), given the differences in HFpEF phenotyping and characteristics between trials
5 and registries. C-statistics were compared to the PREDICT-HFpEF model¹⁹ in EMPEROR-Preserved.
6 In all external validation data model performance was assessed in men and women separately.
7 Additionally, to account for potential geographical differences in average risk and study-specific
8 selection mechanisms, the model was recalibrated using the expected-observed (EO) ratio (a
9 single multiplicative constant) when necessary (see Supplementary Methods for details).

10

11 Sensitivity analyses

12 In EMPEROR-Preserved, model performance was assessed in subgroups of treatment arm
13 (SGLT2 inhibitor or placebo) and stratified by diabetes status and obesity ($BMI \geq 30 \text{ kg/m}^2$). In the
14 NHSESDE, a sensitivity analysis was conducted during the COVID-19 period (follow-up from Jan 1st
15 2020 until March 1st 2022). Detailed description of this COVID cohort can be found in the
16 Supplementary Methods.

17

18 Prediction of individual treatment benefit

19 A potential application of the LIFE-Preserved model is the estimation of individual benefits
20 from starting preventive treatment, as has been done in previous risk models.^{22,34-36} The LIFE-
21 Preserved model can be combined with the latest relative treatment effects from trials and meta-
22 analyses to estimate absolute individual treatment effects, for which a detailed approach is

1 available in Supplementary Methods. The individual treatment benefits can be expressed as
2 absolute risk reduction or gain in HF hospitalization-free life years. If a patient is already receiving
3 one or more evidence-based treatments, the corresponding treatment effects can be combined to
4 LIFE-Preserved in the same way, providing risk estimates that reflect the patient's current
5 treatment regimen. For illustration, the individual treatment benefit of starting a SGLT2 inhibitor
6 (hazard ratio [HR] 0.81)⁴⁰ was estimated for two example patients. Suggested HRs for treatment
7 options are reported in Supplementary Table S5 and serve as illustrative examples derived from
8 current evidence. These should be updated as new treatments and trial results become available.

9 All analyses were conducted using R-statistical programming (Version 4.4.1, R Foundation
10 for Statistical Computing, Vienna, Austria). Results were reported in line with the TRIPOD
11 statement (checklist in Supplementary Material)⁴¹.

12 RESULTS

13 Baseline characteristics

14 The SwedeHF population used for derivation consisted of 10 481 women and 9851 men,
15 with a median age at baseline of 80 years [interquartile range (IQR) 74 – 85] and 77 years [69 – 83],
16 respectively. Detailed sex-specific baseline characteristics are presented in Table 1 and in
17 Supplementary Table S6 for baseline characteristics stratified for in- and outpatients. In women,
18 5044 (48%) first HF hospitalizations or CV deaths occurred during a median follow-up of 1.7 years
19 [interquartile range (IQR) 0.5 – 4.2 years], and 3175 (30%) non-CV deaths occurred over a total
20 median follow-up of 2.7 years [IQR 1.0 – 5.3]. In men, 4297 (44%) first HF hospitalizations or CV
21 deaths occurred during a median follow-up of 1.8 years [IQR 0.6 – 4.2] and 2866 (29%) non-CV
22 deaths occurred over a total median follow-up of 2.6 years [IQR 1.0 – 5.3]. Missingness was high for

1 some predictors [NT-proBNP (45%), BMI (35%), NYHA class (33%), and current smoking (29%)],
2 and were below 6% for other predictors (Supplementary Table S4). Because complete case
3 analysis may lead to loss of statistical power and possible bias, values were imputed using sex-
4 specific multiple imputation by chained equations models.⁴² Assessment of trace plots
5 demonstrated good convergence and stable parameter estimation across the 45 iterations of the
6 multiple imputation models.

8 Model derivation

9 The cause-specific HRs of the LIFE-Preserved model are presented in Table 2. For
10 interpretability, Supplementary Figure S4 illustrates the effect of continuous predictors and age
11 interactions on the primary outcome of HF hospitalization or CV death across the full range of
12 predictor values and ages. All parameters needed for individual predictions are shown in the
13 Supplementary Material; age-specific baseline survivals are presented in Supplementary Table S7
14 and Figure S5, and equations for linear predictors are provided in Supplementary Table S8, detailed
15 explanation is available in Supplementary Methods. Internal validation C-statistic was 0.732 (95%
16 confidence interval [CI] 0.723 – 0.741) for the primary outcome of HF hospitalization or CV death.
17 Discrimination C-statistics were similar in men and women, and higher in outpatients compared to
18 those included from the inpatient setting (Supplementary Table S9). Predicted risks matched
19 observed incidence in internal validation for both outcomes at 2 and 10 years and this alignment
20 was maintained when stratified for in- and outpatients (Supplementary Figure S6).

21

1 External validation

2 External validation of the LIFE-Preserved risk model included data from 41 622 patients with
3 HFpEF in which 15 035 (36%) first HF hospitalizations or CV deaths were observed. Median follow-
4 up times ranged from 1.9 years (IQR 0.7 – 2.8) in the NHSE SDE to 6.8 years (IQR 4.2 – 8.6) in HF-
5 Particles. Detailed patient characteristics of the external validation data sources are presented in
6 Table 3.

7 For the primary outcome of HF hospitalization and CV death pooled C-statistics were 0.714
8 (95% CI 0.652 – 0.775) in trials and 0.658 (0.599 – 0.717) in registries (Figure 1). Model
9 discrimination remained consistent when validated at a prediction horizon of 8 years in HF-
10 Particles (C-statistic 0.693, 95% CI 0.651–0.735) and 10 years in VA (C-statistic 0.687, 95% CI
11 0.642 – 0.732). C-statistics were similar in men and women (Supplementary Figure S7) and higher
12 compared to the existing PREDICT-HFpEF model¹⁹ in EMPEROR-Preserved (Supplementary Table
13 S10). For the competing outcome of non-CV death C-statistics were consistent in external
14 validation (Supplementary Table S9).

15 Predicted risks of HF hospitalization or CV death from the LIFE-Preserved model were in
16 line with the observed incidence in VA, TOPCAT-Americas and HF-Particles (Figure 2). In
17 EMPEROR-Preserved (EO ratio 1.55 for women and 1.31 for men) and the NHSE SDE (EO ratio 1.24
18 for women and 1.28 for men) some overestimation was observed. EO ratios in all external
19 validation cohorts are shown in Supplementary Table S11. Following recalibration, predicted risks
20 of HF hospitalization or CV death were aligned with observed incidence in all validation data
21 sources (Figure 2), and similar in men and women (Supplementary Figure S8-9). The risk of non-CV
22 death was overestimated prior to recalibration in the external trial populations and underestimated
23 in the NHSE SDE (Supplementary Figure S10 and Table S11).

24

1 Sensitivity analyses

2 In EMPEROR-Preserved, performance was similar in subgroups of diabetes status and
3 obesity regarding discrimination (Supplementary Table S9) and calibration (Supplementary Figure
4 S11). After accounting for the treatment effect of the intervention as reported in the EMPEROR-
5 Preserved trial (HR 0.79)⁷, calibration (Supplementary Figure S11) and discrimination
6 (Supplementary Table S9) was similar in the treatment and placebo groups. In a sensitivity analysis
7 of the NHSE SDE comparing the COVID-19 pandemic period with the post-COVID-19 pandemic
8 period (main analysis for the NHSE SDE) similar calibration results were observed (Supplementary
9 Figure S12).

10

11 Prediction of individual treatment benefit

12 A clinical example of the possible use of the LIFE-Preserved model is illustrated in Figure 3.
13 The LIFE-Preserved model can be used to predict the individual 2-year, 10-year and lifetime risk of
14 HF hospitalization or CV death for two example HFPeF patients: a 75-year old woman and a 55-year
15 old man. Combined with HRs from a meta-analysis⁴⁰, individual treatment benefits for starting a
16 SGLT2 inhibitor can be estimated for the two example patients. Risk factor levels of both example
17 patients at baseline are presented in Figure 3. The treatment benefit can be expressed as absolute
18 risk reduction, for example, the reduction in 2- or 10-year risk, or as gain in HF hospitalization-free
19 life years (Figure 3). The gain in HF hospitalization-free life years is 0.6 years for the 75-year old
20 woman, compared with 1.7 years for the 55-year old man. An online interactive calculator for
21 individualized predictions and treatment benefits is available here.

1 DISCUSSION

2 The LIFE-Preserved model for the prediction of short-term and lifetime risks of HF hospitalization or
3 CV death was derived and externally validated using data from 48,394 patients with HFpEF. The
4 LIFE-Preserved model allows the estimation of HF hospitalization-free survival and life expectancy.
5 The model can be combined with the best available evidence from clinical trials to predict
6 individual benefit from guideline-recommended therapies expressed as HF hospitalization-free life
7 years gained. An interactive calculator of the LIFE-Preserved model has been provided until future
8 implementation of the model in online calculators, such as the CE-marked medical device U-
9 prevent.

10 Of the datasets used, SwedeHF was considered the most representative cohort for routine
11 clinical practice, providing real-world incidences of HF hospitalizations, CV and non-CV death.
12 Therefore, the model without any recalibration factors (i.e. based on the event rates in the
13 SwedeHF registry), is recommended for use in daily clinical practice. Predicted risks of HF
14 hospitalization or CV death from the LIFE-Preserved model were in line with the observed incidence
15 in VA, TOPCAT-Americas and HF-Particles. In EMPEROR-Preserved and the NHSE SDE,
16 recalibration of risks was required to reflect observed incidence in those cohorts. This was likely
17 due to the stricter inclusion criteria in clinical trials, differences in patient inclusion and baseline
18 measurement during the COVID period, and/or geographical differences across data sources.
19 Importantly, should new and more broadly representative HFpEF registries become available in the
20 future, the model should be readily recalibrated to reflect local risk levels and increasing use of
21 novel HF therapies. Such recalibration can be easily incorporated into the online implementation,
22 allowing the LIFE-Preserved model to remain accurate and adaptable as new data sources emerge.

23 Regarding clinical implementation, the LIFE-Preserved model is specifically designed to
24 support clinical decision-making by identifying patients at highest risk of a next or first HF

1 hospitalization or CV death. It enables clinicians to estimate prognosis and to guide timely
2 initiation of evidence-based therapies. Importantly, as patient risk profiles evolve over time and
3 after hospitalizations, the model's iterative use allows for updated prognostic assessments and
4 supports ongoing decisions about therapy adjustments, rather than serving as a one-time risk
5 estimate. Consequently, the model can also be applied to patients who have already experienced a
6 HF hospitalization, using their updated clinical profile. The model can be used easily through the
7 online risk calculator and will also be incorporated into the CE-marked medical device U-prevent
8 platform, further facilitating its accessibility across diverse clinical settings.

9 The LIFE-Preserved model has several strengths compared to existing risk models for the
10 HFpEF population. First, LIFE-Preserved can predict risk at any time horizon as well as lifetime risk.
11 Prior risk models are limited to predicting outcomes over a 1- or 2-year period^{15,19}, whereas HFpEF
12 is a chronic disease and therapies are usually continued lifelong. The limited prediction horizon of
13 previously available risk models does not cover the full-time horizon relevant for clinical decision-
14 making. LIFE-Preserved showed good discriminative ability in external validation, which was shown
15 in the current study for up to ten years in external validation data. Pooled C-statistics were higher in
16 trial data than in registry data, likely reflecting more precise phenotyping of HFpEF patients in trials,
17 rather than true differences in model performance. In addition, the higher observed incidence in
18 the registries could also have contributed to a lower a C-statistics in registries.⁴³ In comparison to
19 the existing PREDICT-HFpEF model, the LIFE-Preserved model showed a better discrimination
20 (Supplementary Table S10). LIFE-Preserved also demonstrated adequate calibration in both
21 registry and trial data, making it a useful tool for guiding long-term treatment decisions in this
22 population (Structured Graphical Abstract).

23 Secondly, LIFE-Preserved can estimate lifetime treatment benefits. As a clinical example,
24 the estimated treatment benefit for SGLT2 inhibitors is presented, based on applying clinical trial

1 HRs to modelled absolute risks rather than observed treatment effects in the derivation cohort,
2 where usage was very low. These estimates provide a framework for understanding potential
3 lifetime treatment benefit. The lifetime treatment benefit, defined as the gain in HF hospitalization-
4 free life years, may provide a more intuitive measure for individuals considering preventive
5 treatment. It has been shown that treatment benefits can contribute to risk communication and
6 reduce decisional conflict for individuals.⁴⁴ This information improves shared decision-making,
7 enabling patients and physicians to collaboratively balance expected treatment benefits against
8 practical considerations such as medication burden, potential side effects, and financial costs.

9 Third, the LIFE-Preserved model derives strength from its derivation in SwedeHF, which is
10 one of the largest datasets of HFpEF patients with long-term follow-up for HF hospitalizations and
11 deaths. SwedeHF includes patients from the real-world setting, including both patients from the
12 outpatient setting as well as patients who are hospitalized at inclusion. The LIFE-Preserved model
13 demonstrated good calibration in both groups, indicating that it can be reliably used for stable HF
14 patients as well as for those recently hospitalized. The high observed event rates for HF
15 hospitalization or CV death as well as for non-CV death in SwedeHF are consistent with what has
16 been reported in other real-world HFpEF populations.³ This ensures high generalizability and
17 relevance to real-world patient populations. Risk factor measurements reflect clinical care which
18 has been shown to improve model performance and transferability.⁴⁵ By comparison, trial data
19 usually have more complete baseline data but may be less representative of patients as seen in
20 clinical practice, which could lead to potential misalignment of predicted and observed risk. This is
21 likely due to strict exclusion criteria in trials, which may introduce selection by excluding less
22 healthy individuals. For example, the LIFE-Preserved model, developed in real-world data,
23 overestimated the risk of non-CV death in those trials where both patients with severe

1 comorbidities and limited life expectancy were excluded (TOPCAT-Americas and EMPEROR-
2 Preserved).

3 Fourth, LIFE-Preserved is externally validated in multiple data sources, both trials and
4 observational cohorts, spanning different European and non-European risk regions. Its consistent
5 performance across these settings supports its applicability and reliability in a wide range of
6 healthcare systems and patient populations.

7 Lastly, the model's development and validation is in line with the PROBAST guideline⁴⁶ and
8 TRIPOD statement.⁴¹ It follows established statistical practices, mirroring those used in the LIFE-
9 HF model for HFrEF.²⁵ This results in two complementary models that can predict both short-term
10 and lifetime risk for two major HF subtypes, providing a more holistic tool for clinical decision-
11 making. We recognize that HF with mildly reduced ejection fraction (HFmrEF) was not included in
12 the LIFE-Preserved model, as clinical characteristics and underlying mechanisms more closely
13 resemble HFrEF and predictor–risk relationships may therefore differ from HFpEF.^{47,48} Future
14 studies should evaluate the performance of HFrEF models (such as LIFE-HF) in HFmrEF to
15 complete the continuum of risk prediction for HF patients. Clinical predictors were predefined and
16 selected based on an extensive literature review and a multidisciplinary expert panel, consisting of
17 cardiologists, primary care physicians, epidemiologists, and health services researchers. Selected
18 predictors are readily available in clinical practice, which enhances the model's practical
19 implementation across all lines of care. Furthermore, LIFE-Preserved adjusts for the competing
20 risk of non-CV death. In older and high-risk populations, such as those with heart failure, not
21 accounting for competing risks has been shown to lead to an overestimation of predicted risks,
22 which could result in overtreatment.³³ By accounting for competing risks, the resulting risk
23 estimates better reflect actual patient outcomes. Lastly, sex-specific derivation of coefficients and
24 baseline hazards ensures that LIFE-Preserved captures differences in relative effects of certain

1 predictors and in patterns of risk progression between men and women. External validation
2 showed that model performance was consistent across sexes, confirming that LIFE-Preserved is
3 appropriately calibrated for both men and women in clinical practice.
4

5 Potential limitations also need to be considered. First, registries such as SwedeHF can be
6 selective and have imperfect coverage. SwedeHF captured 31% of prevalent HF diagnoses in
7 2020.^{26,49} Although this may introduce some degree of selection, the registry nevertheless reflects a
8 broad and clinically relevant HF population. Importantly, the strong and consistent performance of
9 our model in the external validation cohorts provides reassurance that any potential limitations in
10 representativeness did not materially affect model development or applicability. Second, due to
11 the use of routinely collected data, there was a relevant proportion of missing data for several
12 model predictor variables such as NT-proBNP, BMI and NYHA class. However, missing data were
13 imputed using recommended multiple imputation methods, and the model was derived by pooling
14 the coefficients across all 45 imputed datasets. The model performance in external validation
15 supports the robustness of these derived coefficients. Third, although the median follow-up of 1.8
16 years may appear short, this primarily reflects the rapid occurrence of events rather than
17 inadequate observation time. The large population provided substantial long-term data, with
18 follow-up exceeding 4.2 years in 25% (N = 5083) of patients (Supplementary Figure S13a), ensuring
19 sufficient overlapping data across age groups for reliable lifetime risk estimation. Consequently,
20 survival curves remained stable well beyond 10 years (Supplementary Figure S13b). Although true
21 lifetime risk estimates cannot be validated, because no contemporary cohort has sufficiently long
22 follow-up, the model demonstrated good calibration up to 10 years of follow-up in external
23 validation and the methodology used has been shown to yield accurate risks up to at least 17
24 years.³⁴ Fourth, in the current examples, individual treatment benefits were shown only for the

1 outcome of HF hospitalization or CV death, given its relevance and modifiability in HFpEF.
2 Recommendations for these therapies are presently based on observed reductions in HF
3 hospitalizations and CV mortality rather than total mortality. In addition, individual treatment
4 benefits were estimated using a fixed treatment effect (i.e. a single treatment effect regardless of
5 patient characteristics). This approach assumes no heterogeneity in relative treatment response
6 among HFpEF patients, as trials and meta-analyses to date have not reported major differences.
7 However, when applying the model in clinical practice to calculate treatment effects, it should be
8 used alongside the most up-to-date evidence, including any potential heterogeneity in treatment
9 response, should such evidence emerge. The LIFE-Preserved model is designed to provide a
10 framework to be used alongside the latest evidence on therapy effectiveness in this group. The
11 online implementation will be continuously updated, integrating new treatments and updating HRs
12 when new evidence emerges.

13
14 In conclusion, the LIFE-Preserved model can predict risk at any time horizon as well as
15 lifetime risk of HF hospitalization or CV death and HF hospitalization-free life expectancy for
16 individual patients with HFpEF. Combining the LIFE-Preserved model with treatment effects from
17 trials or meta-analyses allows for individualized estimates of treatment benefit. Thus, this model
18 could serve as a tool to improve the management of patients with HFpEF by facilitating
19 personalized medicine and shared decision-making.

21 ESC Cardiovascular Risk Collaboration

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6 Acknowledgements

7 The authors thank all staff members at all care units in Sweden for their contribution to the
8 Swedish HF registry. The authors also acknowledge the contribution of the EMPEROR-
9 Preserved, TOPCAT and HF-Particles investigators. This study was partly performed using
10 data and resources available in the Department of Veterans Affairs Cleveland VA
11 Medical Center. The results and opinions expressed in this manuscript are those of the
12 authors and do not represent those of the Department of Veterans Affairs or the United
13 States. Analysis of VA data was performed by SVD, a full-time employee and researcher in
14 the Department of Veterans Affairs Cleveland VA Medical Center and he takes
15 responsibility for those analyses. All other authors did not have access to the data and
16 received only the results for review.

17 We gratefully acknowledge the contribution of the ESC Cardiovascular Risk Collaboration:
18 Prof. Emanuele Angelantonio (Department of Public Health and Primary Care, University of
19 Cambridge, Victor Phillip Dahdaleh Heart & Lung Research Institute), Prof. Ana Abreu
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24 (Department of Public Health and Primary Care, University of Cambridge, Victor Phillip
25 Dahdaleh Heart & Lung Research Institute). This paper is based on research using data
26 from data contributors of the EMPEROR-Preserved trial that has been made available
27 through Vivli, Inc. Vivli has not contributed to or approved, and is not in any way
28 responsible for, the contents of this publication. This work was carried out with the support
29 of the BHF Data Science Centre led by Health Data Research UK (BHF Grant no.
30 SP/19/3/34678). This study made use of anonymised data held in NHS England's Secure
31 Data Environment service for England and made available via the BHF Data Science
32 Centre's CVD-COVID-UK/COVID-IMPACT consortium. This work used data provided by
33 patients and collected by the NHS as part of their care and support. We would also like to
34 acknowledge all data providers who make health relevant data available for research. The
35 BHF Data Science Centre's Health Data Science Team provided data curation resources
36 and support.

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- 29

1 Legends

2 Graphical abstract

3 Graphical abstract of the derivation, validation and clinical utility of the LIFE-Preserved model. For visual clarity, the
4 clinical example illustrates a subset of predictors but LIFE-Preserved is based on 14 routinely available predictors: age,
5 sex, diabetes mellitus, atrial fibrillation, current smoking, prior hospitalization for heart failure (HF), chronic obstructive
6 pulmonary disease (COPD), New York Heart Association (NYHA) class, ischaemic heart disease, N-terminal pro-B-type
7 natriuretic peptide (NT-proBNP), body mass index (BMI), heart rate, haemoglobin, and estimated glomerular filtration rate
8 (eGFR). CV = cardiovascular; CVD = cardiovascular death; HFH = heart failure hospitalization; HFpEF = heart failure with
9 preserved ejection fraction; SGLT2i = sodium–glucose cotransporter 2 inhibitor. Created in <https://BioRender.com>

10

11 Figure 1: C-statistics for HF hospitalization or CV death in external validation data

12 Forest plot displaying the discrimination in all external validation data is based on Harrell's C-statistic for HF
13 hospitalization or CV death adjusted for non-CV death. Discrimination was assessed at the 75th percentile of follow-up
14 duration (2-year risks for EMPEROR-Preserved and the NHSE SDE, 4-year risks for TOPCAT-Americas, 8-year risks for HF-
15 Particles) and at 10-year risks for VA. Pooled C-statistics were calculated using inverse-variance weighting. Cohort sizes
16 here may be lower than the total cohort population listed in Table 3 as risks can only be estimated up to age 90 (e.g., 89-
17 year-olds can have 1-year risk but no risks for 2-year or higher and are consequently excluded at the horizon for
18 validation). CI = confidence interval.

19

20 Figure 2: Calibration for HF hospitalization or CV death in external validation data

21 Calibration of the LIFE-Preserved model in external validation. Risks were recalibrated in the NHSE SDE and EMPEROR-
22 Preserved using EO ratios specified in Supplementary Table S11. Smoothed calibration plots of predicted risks versus
23 observed risks and a histogram of the predicted risks (in blue) is shown. Cohort sizes here may be lower than the total
24 cohort population listed in Table 2 as risks can only be estimated up to age 90 (e.g. 89-year-olds can have 1-year risk but
25 no risks for 2-year or higher and are consequently excluded at the horizon for validation). HF = heart failure; CV =
26 cardiovascular.

27 Figure 3: Clinical example of estimation of individual treatment benefit

28 Clinical example of estimation of individual short-term and lifetime risk and individual treatment benefit after starting a
29 SGLT2 inhibitor for two different HFpEF patients. Event-free survival in the curves is defined as survival without any HF
30 hospitalization or CV death. Treatment benefits were estimated by combining the individual absolute risks from the LIFE-
31 Preserved model with the published hazard ratios from trials or a meta-analyses. In this example the published hazard
32 ratio of 0.81 was used as treatment effect of starting a SGLT2 inhibitor.⁴⁰ Treatment benefits do not reflect observed
33 treatment use in the cohorts and should be interpreted as projections of potential benefit. The individual treatment
34 benefit can be shown as an absolute risk reduction (barchart, bottom row) or as gain in HF-hospitalization free lifeyears
35 (dark green area in survival plot, top row).

36 HFpEF = heart failure with preserved ejection fraction; NYHA = New York Heart Association; BMI = body mass index; bpm
37 = beats per minute; NT-proBNP = N-terminal pro-B-type natriuretic peptide; eGFR = estimated glomerular filtration rate;
38 Hb = haemoglobin; COPD = chronic obstructive pulmonary disease; HF = heart failure; CV = cardiovascular; SGLT2 =
39 sodium–glucose cotransporter 2. Created in <https://BioRender.com>

40

	Women (n = 10 481)	Men (n = 9851)
Age (years)	80 [74–85]	77 [69–83]
Current smokers	867 (8)	891 (9)
Physical signs		
Body mass index (kg/m ²)	27 [23–32]	27 [24–31]
Systolic blood pressure (mmHg)	130 [120–148]	130 [120–145]
Diastolic blood pressure (mmHg)	70 [65–80]	73 [65–80]
Heart rate (bpm)	72 [64–83]	70 [61–80]
Clinical features of HF		
NYHA class		
I	1390 (13)	1685 (17)
II	4809 (46)	4549 (46)
III	3897 (37)	3291 (33)
IV	385 (3.7)	326 (3.3)
Duration of HF (years)	0.2 [0.0–2.6]	0.2 [0.0–2.8]
Left bundle branch block	620 (6)	732 (7)
Medical history		
Prior HFH	5833 (56)	5388 (55)
Recent HFH (≤ 6 months)	4829 (46)	4202 (43)
Diabetes mellitus	2752 (26)	2989 (30)
Ischaemic heart disease	4494 (43)	4866 (49)
COPD	1739 (17)	1572 (16)
Obesity ^a	3492 (33)	3032 (31)
Atrial fibrillation	6665 (64)	6435 (65)
Stroke or transient ischaemic attack	1875 (18)	1809 (18)
Peripheral artery disease	858 (8)	1046 (11)
Valve disease	3662 (35)	3136 (32)
Dilated cardiomyopathy	271 (3)	455 (5)
Laboratory measurements		
NT-proBNP (pg/mL)	2120 [972–4343]	1755 [722–3700]
Haemoglobin (mmol/L)	7.8 [7.1–8.4]	8.2 [7.3–8.9]
Estimated GFR (mL/min/1.73 m ²)	55 [40–71]	46 [33–59]
Medication use		
Loop diuretics	8552 (82)	7578 (77)
ACE-inhibitor/ARB	7404 (71)	7364 (75)
Beta-blocker	8583 (82)	7897 (80)
Statin	4010 (38)	4779 (48)
MRA	3164 (30)	2937 (30)
Digoxin	1739 (17)	1125 (11)
Nitrate	1624 (16)	1333 (14)
Device therapy ^b	134 (1)	266 (3)
ARNI	24 (0.2)	44 (0.4)
SGLT2 inhibitor	25 (0.2)	68 (0.7)
GLP-1 receptor agonist	44 (0.4)	79 (0.8)
Outcomes: incidence (%) event rate per 1000 PY		
First HFH/CV death	5044 (48) 169	4297 (44) 148
First HFH	3735 (36) 125	3226 (33) 111
Total number of HFH	9158 (36) 307	8575 (33) 296
Non-CV death	3175 (30) 83	2866 (29) 80
CV death	2776 (26) 73	2293 (23) 64
Follow-up time for first HFH (years)	1.7 [0.5–4.2]	1.8 [0.6–4.2]

Follow-up time for death (years)

2.7 [1.0–5.3]

2.6 [1.0–5.3]

1 **Table 1.** Sex-specific baseline characteristics of SwedeHF

2 Data are presented as *n* (%) or median [25th–75th percentile].

3 HF = heart failure; NYHA = New York Heart Association; COPD = chronic obstructive pulmonary disease; NT-proBNP = N-
4 terminal pro-B-type natriuretic peptide; GFR = glomerular filtration rate (calculated with the 2009 Chronic Kidney Disease
5 Epidemiology Collaboration formula); ACE = angiotensin-converting enzyme; ARB = angiotensin receptor blocker; ARNI =
6 angiotensin receptor–neprilysin inhibitor; MRA = mineralocorticoid receptor antagonist; SGLT2 = sodium-glucose
7 cotransporter 2; GLP-1 = glucagon-like peptide-1; HFH = heart failure hospitalization; CV = cardiovascular; PY = patient-
8 years.

9 ^a Obesity is defined as body mass index ≥ 30 kg/m².

10 ^b Device therapy includes cardiac resynchronization therapy and implantable cardioverter-defibrillator.

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ACCEPTED MANUSCRIPT

1

	Hazard ratios (95% CI)			
	HF hospitalization or CV death		Non-CV death	
	Women	Men	Women	Men
Age at baseline (per 10 years)	1.61 (1.38–1.89)	1.58 (1.35–1.86)	0.73 (0.63–0.86)	0.71 (0.60–0.83)
Current smoking (vs never/former)	1.16 (1.02–1.32)	1.19 (1.04–1.36)	0.96 (0.82–1.12)	1.30 (1.11–1.52)
NYHA class III/IV (vs I/II)	1.24 (1.16–1.34)	1.34 (1.24–1.44)	1.37 (1.26–1.50)	1.29 (1.17–1.43)
Ischaemic heart disease	1.20 (1.13–1.27)	1.15 (1.08–1.23)	0.86 (0.80–0.93)	0.86 (0.80–0.93)
COPD	1.30 (1.20–1.41)	1.33 (1.22–1.44)	2.05 (1.88–2.25)	1.76 (1.60–1.93)
Atrial fibrillation	1.21 (1.13–1.29)	1.16 (1.08–1.25)	0.86 (0.79–0.93)	0.90 (0.82–0.98)
Prior HF hospitalization (at any time)	1.38 (1.28–1.48)	1.38 (1.28–1.49)	1.14 (1.06–1.24)	1.23 (1.13–1.33)
HF hospitalization in past 6 months ^a	2.06 (1.91–2.23)	1.97 (1.81–2.14)	1.26 (1.13–1.41)	1.26 (1.12–1.41)
Diabetes mellitus	1.53 (1.38–1.69)*	1.39 (1.28–1.52)*	1.35 (1.32–1.38)	1.26 (1.23–1.28)
NT-proBNP (4000 vs 800 pg/mL)	1.47 (1.38–1.57)	1.50 (1.40–1.61)	1.44 (1.33–1.57)	1.33 (1.22–1.46)
BMI (35 vs 25 kg/m ²)	1.14 (1.05–1.23)	1.24 (1.11–1.38)	0.88 (0.80–0.98)	0.94 (0.82–1.06)
Haemoglobin (7 vs 8.5 mmol/L)	1.10 (1.04–1.15)	1.13 (1.08–1.19)	1.21 (1.14–1.28)	1.28 (1.20–1.35)
Estimated GFR (35 vs 65 mL/min/1.73 m ²)	1.24 (1.17–1.31)	1.17 (1.10–1.25)	1.29 (1.21–1.38)	1.12 (1.04–1.20)
Heart rate (80 vs 60 bpm)	1.06 (1.00–1.12)	1.14 (1.08–1.21)	1.22 (1.13–1.31)	1.18 (1.11–1.27)

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3 **Table 2.** Hazard ratios of the predictors from the LIFE-Preserved model

4 The presented hazard ratios are equal to the exponential of the coefficients and they cannot be causally interpreted. Hazard ratios for continuous predictors are shown
5 for clinically interpretable comparisons, for insight in the detailed relationship of the continuous predictors and age interactions see Figure S4.

6 CI = confidence interval; HR = hazard ratio; NYHA = New York Heart Association; COPD = chronic obstructive pulmonary disease; HF = heart failure; NT-proBNP = N-
7 terminal pro-B-type natriuretic peptide; BMI = body mass index; eGFR = estimated glomerular filtration rate (calculated with the 2009 Chronic Kidney Disease
8 Epidemiology Collaboration formula).

9 * Interaction with age, hazard ratio shown for age 70 years.

10 ^aTime-varying effect, only applies to the first year of risk predictions and is on top of the effect of prior HF hospitalization at any time.

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	Trials		Registries		
	EMPEROR- Preserved (n = 3955)	TOPCAT-Americas (n = 1553)	NHSE SDE (n = 17 005)	HF-Particles (n = 907)	VA (n = 4642)
Age (years)	74 [67–79]	73 [64–79]	79 [72–85]	71 [63–77]	66 [62–71]
Women	1988 (50)	802 (52)	9290 (55)	457 (50)	62 (1)
Current smokers	248 (6)	90 (6)	1035 (6)	16 (2)	935 (20)
Physical signs					
Body mass index (kg/m ²)	30 [26–34]	33 [28–39]	30 [26–35]	28 [25–32]	31 [28–35]
Heart rate (bpm)	69 [61–78]	68 [61–76]	74 [65–84]	78 [70–84]	*
Clinical features of HF					
NYHA class (III/IV)	736 (19)	556 (36)	12 190 (72)	355 (39)	1727 (37)
Medical history					
Prior HFH	871 (22)	899 (58)	17 005 (100)	82 (9)	983 (21)
Recent HFH (<6 months)	871 (22)	239 (15)	7145 (42)	67 (7)	769 (17)
Diabetes mellitus	1896 (48)	692 (45)	7915 (47)	266 (29)	2447 (53)
Ischaemic heart disease	1125 (28)	699 (45)	10 575 (62)	359 (40)	4642 (100)
COPD	537 (14)	252 (16)	5740 (34)	43 (5)	1101 (24)
Atrial fibrillation	2147 (54)	643 (41)	11 525 (68)	311 (34)	358 (8)
Laboratory measurements					
NT-proBNP (pg/mL)	941 [478–1672]	738 [432–1393]	3717 [1549–6749]	*	*
Haemoglobin (mmol/L)	8.2 [7.6–8.8]	7.9 [7.3–8.7]	7.3 [6.4–8.1]	8.2 [7.6–8.8]	8.4 [7.6–9.1]
Estimated GFR (mL/min/1.73 m ²)	59 [45–74]	58 [47–73]	51 [37–68]	83 [65–93]	73 [58–90]
Follow-up time (years)	2.0 [1.4–2.7]	2.5 [1.4–3.9]	1.9 [0.6–2.8]	6.8 [4.2–8.6]	3.3 [0.9–6.3]
Incidence of HFH/CV death	579 (15)	451 (29)	6810 (40)	341 (38)	1749 (38)
Event rate of HFH/CV death per 1000 PY	73	110	241	60	98
Event rate of non-CV death per 1000 PY	25	29	137	19	38

1 **Table 3.** Baseline characteristics in external validation data

2

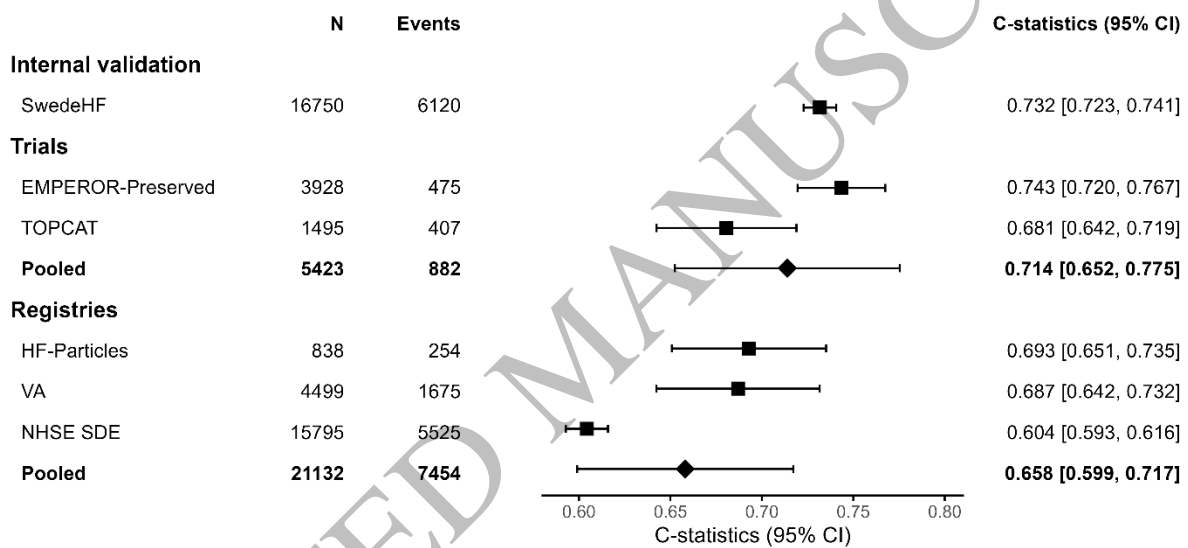
3 Data are presented as *n* (%) or median [25th–75th percentile].

4 HF = heart failure; NYHA = New York Heart Association; COPD = chronic obstructive pulmonary disease; NT-proBNP = N-
 5 terminal pro-B-type natriuretic peptide; eGFR = estimated glomerular filtration rate (calculated with the 2009 Chronic
 6 Kidney Disease Epidemiology Collaboration formula); HFH = heart failure hospitalization; CV = cardiovascular; PY =
 7 patient-years.

8 * Data for this variable is missing completely in this data source.

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Figure 1
 165x75 mm (DPI)

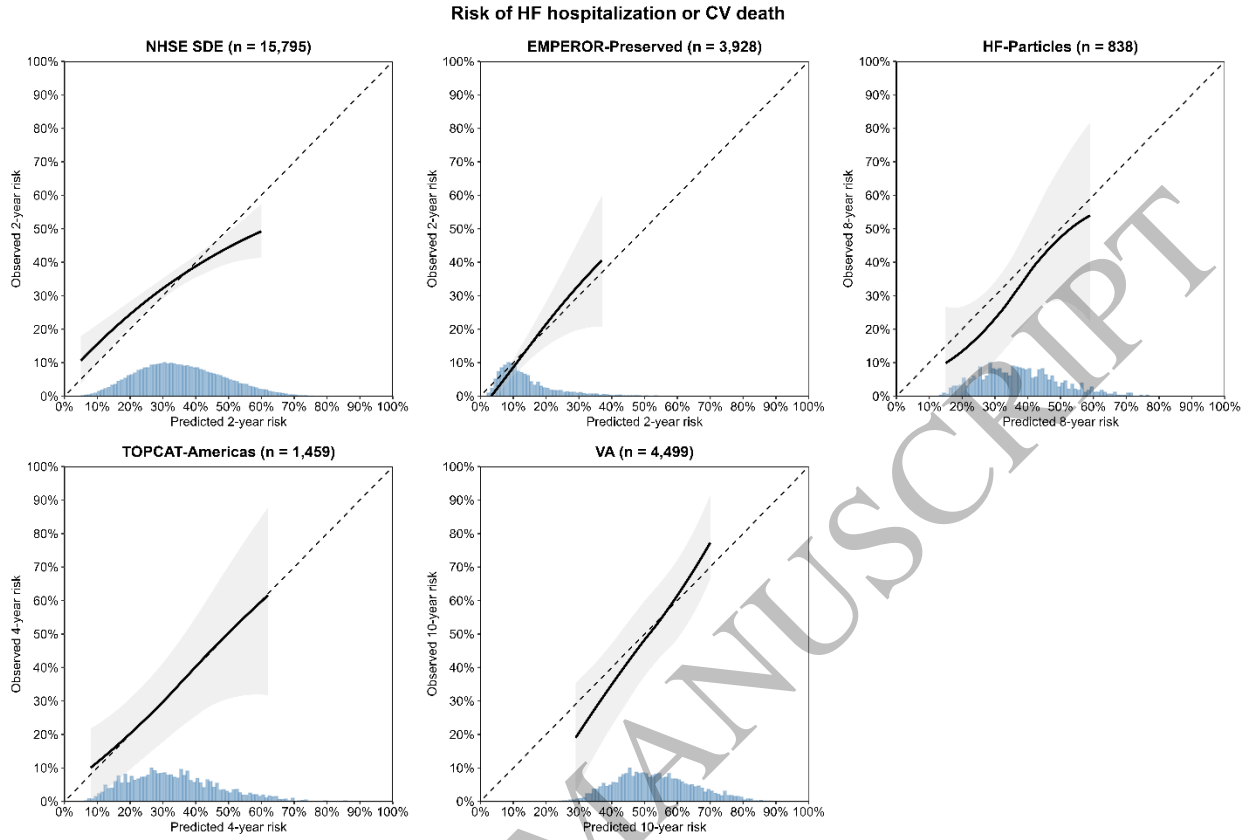
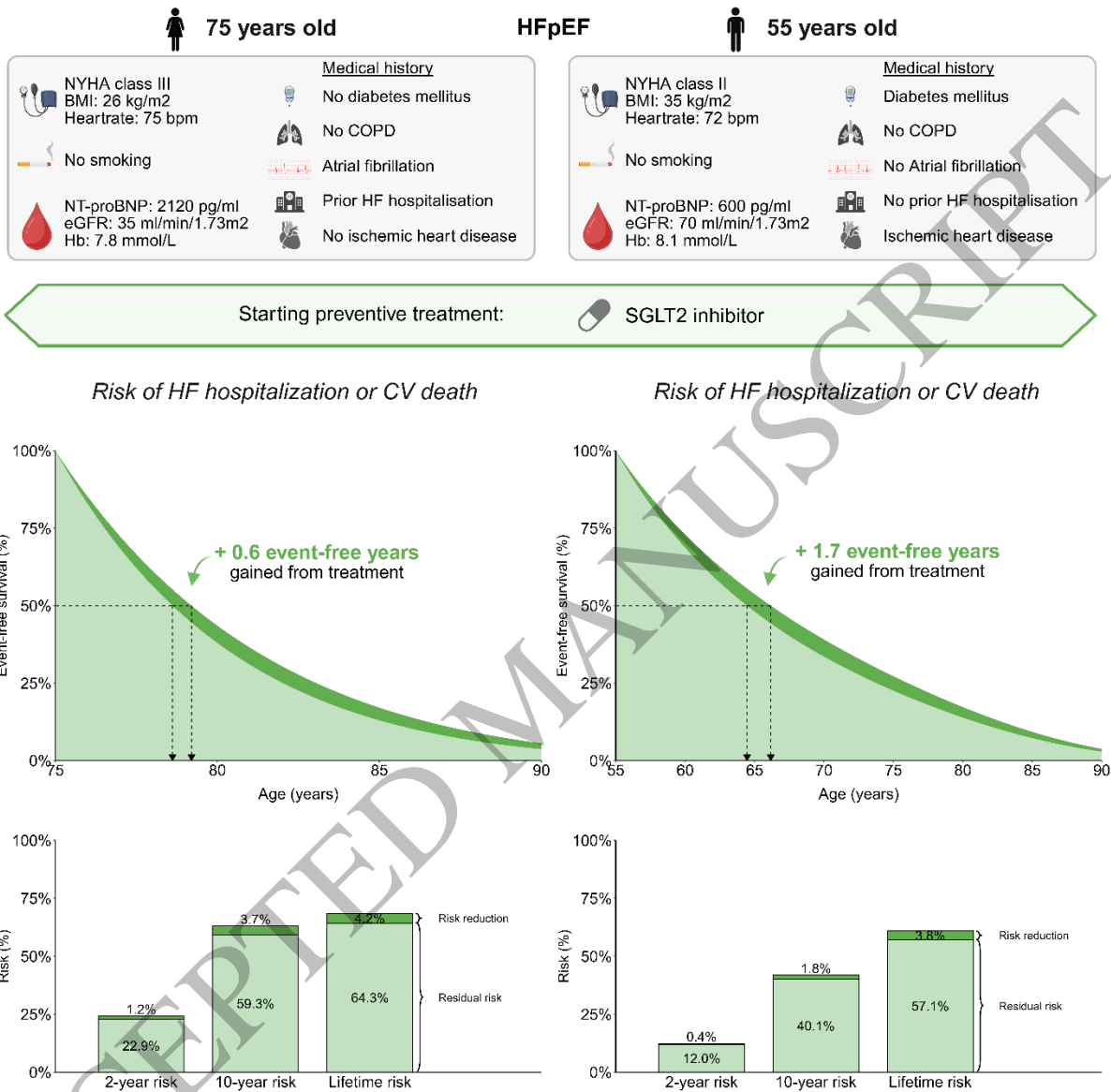


Figure 2
165x110 mm (DPI)

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Figure 3
165x179 mm (DPI)

Key Question

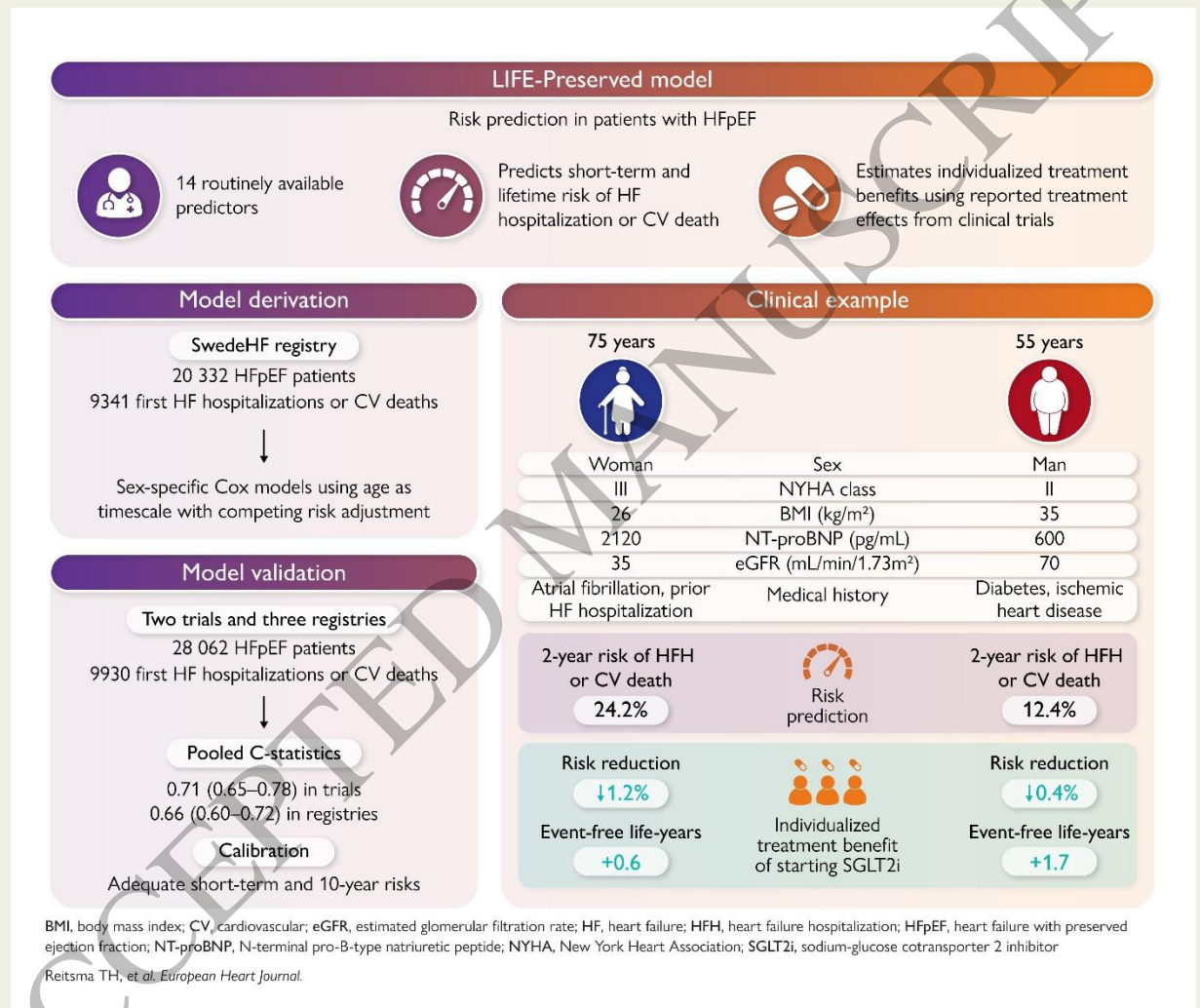
How can patients with heart failure with preserved ejection fraction (HFpEF) at risk of heart failure (HF) hospitalization or cardiovascular death be identified in clinical practice using easily available clinical information?

Key Finding

The LIFE-Preserved model showed good discrimination and calibration for HF hospitalization or cardiovascular death in HFpEF across multiple international trials and real-world registries, with comparable performance in women and men.

Take Home Message

The LIFE-Preserved score may help identify high-risk HFpEF patients who may benefit most from preventive strategies.



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Graphical Abstract
165x178 mm (DPI)