



Risk Prediction Models for Mortality in Ambulatory Patients With Heart Failure: A Systematic Review

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Original Article

Risk Prediction Models for Mortality in Ambulatory Patients With Heart Failure

A Systematic Review

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Background—Optimal management of heart failure requires accurate assessment of prognosis. Many prognostic models are available. Our objective was to identify studies that evaluate the use of risk prediction models for mortality in ambulatory patients with heart failure and describe their performance and clinical applicability.

Methods and Results—We searched for studies in Medline, Embase, and CINAHL in May 2012. Two reviewers selected citations including patients with heart failure and reporting on model performance in derivation or validation cohorts. We abstracted data related to population, outcomes, study quality, model discrimination, and calibration. Of the 9952 studies reviewed, we included 34 studies testing 20 models. Only 5 models were validated in independent cohorts: the Heart Failure Survival Score, the Seattle Heart Failure Model, the PACE (incorporating peripheral vascular disease, age, creatinine, and ejection fraction) risk score, a model by Frankenstein et al, and the SHOCKED predictors. The Heart Failure Survival Score was validated in 8 cohorts (2240 patients), showing poor-to-modest discrimination (*c*-statistic, 0.56–0.79), being lower in more recent cohorts. The Seattle Heart Failure Model was validated in 14 cohorts (16057 patients), describing poor-to-acceptable discrimination (0.63–0.81), remaining relatively stable over time. Both models reported adequate calibration, although overestimating survival in specific populations. The other 3 models were validated in a cohort each, reporting poor-to-modest discrimination (0.66–0.74). Among the remaining 15 models, 6 were validated by bootstrapping (*c*-statistic, 0.74–0.85); the rest were not validated.

Conclusions—Externally validated heart failure models showed inconsistent performance. The Heart Failure Survival Score and Seattle Heart Failure Model demonstrated modest discrimination and questionable calibration. A new model derived from contemporary patient cohorts may be required for improved prognostic performance. (Circ Heart Fail. 2013;6:881-889.)

Key Words: heart failure ■ prediction models ■ prognosis ■ survival

Heart failure (HF) is a frequent health problem with high morbidity and mortality, increasing prevalence and escalating healthcare costs. ^{1,2} Older patient age, multiple comorbidities, and different patterns of disease progression create important challenges in patient management. Because the impact of these factors and their interactions remain incompletely understood, predicting patients' clinical course is difficult.

Editorial see p 877 Clinical Perspective on p 889

Accurate estimation of prognosis is important for many reasons. Patients are concerned about their probability of future events. Physicians may use prognosis estimates to decide the appropriate type and timing of additional tests or therapies,

including heart transplantation and mechanical circulatory support. Accurate prognostic assessment may prevent delays in appropriate treatment of high-risk patients or overtreatment of low-risk patients. Knowledge of prognosis also facilitates research, for instance in the design of randomized trials and the exploration of subgroup effects.

To be usefully applied, prognostic models must be accurate and generalizable. Models may be inaccurate because of omission of important predictors, derivation from unrepresentative cohorts, overfitting or violations of model assumptions.

In the past 3 decades, investigators have developed many models to predict adverse outcomes in patients with HF.^{3,4} Clinicians and researchers wishing to use prognostic models would benefit from knowledge of their characteristics and

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performance. Therefore, we performed a systematic review to identify studies evaluating the use of risk prediction models for mortality in ambulatory patients with HF and to describe their performance and their clinical applicability.

Methods

Data Sources and Searches

In May 2012, with the assistance of an experienced research librarian, we performed a systematic search of electronic databases, including Medline, Embase, and CINAHL. We used several related terms: (internal cardiac defibrillator [ICD]), (heart or cardiac), (mortality or survival), and (multivariate analysis or regression analysis or risk factor or prediction or prognostic factor). The full search strategy is outlined in Appendix A in the online-only Data Supplement (Methods in the online-only Data Supplement). We identified additional studies by searching bibliographic references of included publications.

Study Selection

Eligible articles enrolled adults (>19 years) who were ambulatory patients with HF; used multivariable analysis (≥2 independent variables) to predict mortality or a composite outcome including mortality; reported >30 deaths; reported results as a score, a prediction rule, or as a set of regression coefficients sufficient to make predictions for individual patients; and reported a measure of discrimination or calibration. We also included studies evaluating the performance of an existing score in a different population to the one from which it was developed, and reported model discrimination and calibration. There were no restrictions on study design, left ventricular ejection fraction (LVEF), language, or date of publication. We excluded studies that enrolled patients during hospital admission or duplicate studies providing no new relevant data.

Two reviewers independently screened titles and abstracts, and then evaluated full-text versions of all articles deemed potentially relevant by either reviewer. During full-text screening, in cases of disagreement, consensus was reached through discussion. If consensus could not be reached, a third reviewer resolved the issue. Agreement between reviewers was assessed using weighted κ (0.92). Appendix B in the online-only Data Supplement (Methods in the online-only Data Supplement) shows the eligibility form.

Data Extraction

From each study, we abstracted data related to eligibility criteria, data source, time frame of recruitment, and characteristics of the population, including age, sex, ischemic cardiomyopathy, LVEF, use of β-blockers and ICD, definition, and number of events. We also identified variables included in the prediction models.

Assessment of Study Quality, Model Adequacy, and Performance

The assessment of study quality and model performance was based on what authors reported in their published articles. The selection of items for the assessment of study quality, model adequacy, and performance was based on the criteria proposed by Concato et al⁵ and Moons et al.6 Items included whether patient selection was consecutive, whether the data were collected prospectively, whether the percentage of missing data were small (<5%) and was correctly managed (ie, using data imputation), whether patients lost to follow-up were infrequent (<1%), and whether predictors were coded clearly.

To assess model adequacy, we abstracted information related to model derivation, including selection of the variables, coding, linearity of the response for continuous variables, overfitting,7 and model assumptions. To assess model performance, we abstracted data related to discrimination and calibration. Discrimination expresses the extent to which the model is capable of differentiating patients who had events from those who did not. It is commonly assessed using the cstatistic, which is equivalent to the area under the receiver-operating characteristic curve.8 Model discrimination was deemed as poor if

the c-statistic was between 0.50 and 0.70, modest between 0.70 and 0.80, and acceptable if >0.80.9 To assess how changes in HF treatment might modify model performance, we evaluated the impact of β-blockers, use of ICD, and study recruitment date on model discrimination graphically including models tested in >1 external cohort.

The calibration and goodness-of-fit of a model involves investigating how close the values predicted by the model are to the observed values. We identified the method used to assess model calibration (ie, Hosmer-Lemeshow test or deviance, Cox-Snell analysis, correlation between observed versus predicted events) and estimate of performance.

Table I in the online-only Data Supplement explains the criteria used to assess model adequacy and performance in more detail. Items that were not relevant (eg, in studies validating a preexisting model) were coded as nonapplicable.

Data Synthesis

We summarized the data, focusing on the characteristics of the population from whence models were derived and validated, and the models' performance. We report findings in 2 sections according to external validation (models that were or were not validated in an independent cohort were summarized separately).

Results

After duplicate citations were removed, we screened 6917 citations and ultimately selected 32 studies evaluating 20 prediction models (Figure 1). Only 5 of these models 10-14 were validated in an independent cohort. Among the remaining 15 models, 6 were internally validated by bootstrap; the remaining models were not validated.

Prediction Models Validated in an **Independent Cohort**

The Heart Failure Survival Score (HFSS), 10 the Seattle Heart Failure Model (SHFM),¹¹ the model proposed by Frankenstein et al, 12 the PACE risk score, 13 and the SHOCKED predictors 14 were validated in a different cohort of patients with HF from the model derivation cohort. Tables II and III in the onlineonly Data Supplement, and the Table summarize the characteristics of studies included, the assessment of study quality and model characteristics, respectively.

Heart Failure Survival Score

The HFSS includes 7 variables to predict a composite outcome of death, urgent (UNOS [United Network for Organ Sharing] status 1) heart transplantation and ventricular assist device implantation. Two predictors are binary: ischemic cardiomyopathy and presence of intraventricular conduction delay (QRS >120 ms); and 5 are continuous: LVEF, resting heart rate, mean blood pressure, peak oxygen consumption, and serum sodium. Scores are then divided into 3 categories: high risk, medium risk, and low risk according to prespecified thresholds. 10 The HFSS was derived from a single center cohort including 268 patients with HF and has been validated in 8 independent single-center cohorts including a total of 2240 HF patients. 10,14–19

The validation cohorts involve a broad variety of patient populations (Table II in the online-only Data Supplement), with a mean age from 51 to 70 years, mostly males (65%-82%) with a mean LVEF between 20% and 30%. In 3 cohorts, the frequency of use of β -blockers was <30% and in the remaining 4 cohorts was 64% to 80%. In 4 studies reporting ICD status, the frequency of ICD use was 11%, 19%, 49%, and 78%.

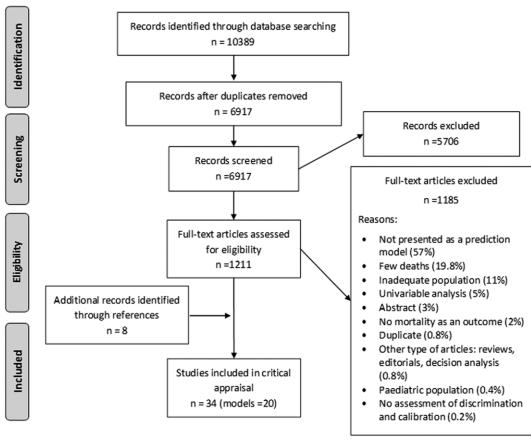


Figure 1. Study selection process. Number of studies during selection.

Model discrimination (assessed by the c-statistic at 1 year) in validation cohorts ranged from poor to modest (0.56–0.79), being modest (between 0.70 and 0.79) in 6 (75%) of the 8 validation cohorts. As shown in Figure 2, model discrimination was worse in cohorts with more frequent use of β -blockers or ICDs, and in more recent studies. Discrimination was poor

(*c*-statistic, <0.70) in validation cohorts in which the rate of ICD use was >40%, studies with a contemporary recruitment date and in 3 of 4 cohorts in which the use β -blockers was >60%. The study by Zugck et al¹⁵ reported a substantially higher discrimination (*c*-statistic=0.84 at 1 year) when peak oxygen consumption was replaced by the 6-minute walk test.

Table. Model Derivation and Performance

Study	Derivation Validation	Model/Variables	Selection	Linear Gradient	Overfitting	Model Assumptions	Calibration	Discrimination (<i>c</i> -Statistic)
Aaronson et al ¹⁰	Derivation	HFSS: Heart rate BP LVEF Sodium Ischemic CMP IVCD Peak Vo ₂	Based on univariable analysis	n.r.	Yes (109 events and 11 variables)	Held	n.r.	At 1 y=0.79 (0.76-0.82)
	Validation in a different cohort	HFSS	n/a	n/a	n/a	n.r.	n.r.	At 1 y=0.76 (0.72–0.80) Overall=0.69 (0.62–0.76)
Zugck et al15	Validation	HFSS	n/a	n/a	n/a	n.r.	n.r.	Overall=0.74 (0.70-0.78)
		HFSS replacing peak VO2 by 6'WT	n/a	n/a	No	n.r.	n.r.	Overall=0.83 (0.79-0.87)
Koelling et al ¹⁶	Validation	HFSS	n/a	n/a	n/a	n.r.	n.r.	Not β -blockers: at 1 y=0.76 (0.72–0.80) β -Blockers: at 1 y=0.73 (0.68–0.78)
Parikh et al ¹⁷	Validation	HFSS	n/a	n/a	n/a	n.r.	n.r.	At 1 y=0.76 (0.70–0.83) (<i>Continued</i>)

Table. Continued

Study	Derivation Validation	Model/Variables	Selection	Linear Gradient	Overfitting	Model Assumptions	Calibration	Discrimination (<i>c</i> -Statistic)
Gorodeski et al ¹⁸	Validation	HFSS	n/a	n/a	n/a	n/a	Tested graphically: overestimated survival in HT candidates and more pronouncedly in non-HT candidates	At 1 y: In HT candidates=0.53 (0.50-0.63) In non-HT candidates=0.62 (0.55-0.68)
Goda et al ¹⁹⁻²¹	Validation	HFSS	n/a	n/a	n/a	n.r.	n.r	*At 1 y: Total cohort=0.72 (0.67–0.76) European American (n=417) =0.69 (0.63–0.75) Black (n=125) =0.73 (0.63–0.84) Hispanic American (n=123) =0.76 (0.66–0.85) ICD/CRT patients (n=382) =0.69 (0.63–0.75)
Levy et al ¹¹	Derivation	SHFM: Sex Age NYHA Sodium Uric acid Cholesterol Hemoglobin Lymphocytes Systolic BP LVEF Ischemic CMP Statin Allopurinol Diuretic dose β-blockers ACEI ARB K-sparing diuretic	Based on univariable analysis, forward elimination effect of some treatments were obtained from previous RCTs or meta- analysis	Checked	No	n.r.	Assessed graphically observed vs predicted survival by deciles and by correlation (<i>r</i> =0.97)	
	Validation ELITE2	SHFM	n/a	n/a	n/a	n/a	Correlation (<i>r</i> =0.97)	At 1 y=0.67 (0.65–0.71)
	Validation RENAISSANCE	SHFM	n/a	n/a	n/a	n/a	Correlation (<i>r</i> =0.97)	At 1 y=0.69 (0.68–0.72)
	Validation Val-HeFT	SHFM	n/a	n/a	n/a	n/a	Correlation (<i>r</i> =0.98)	At 1 y=0.81 (0.72-0.90)
	Validation IN-CHF	SHFM	n/a	n/a	n/a	n/a	Correlation (<i>r</i> =0.99)	At 1 y=0.75 (0.70-0.80)
	Validation UW	SHFM	n/a	n/a	n/a	n/a	Correlation (r=0.99)	At 1 y=0.68 (0.63-0.73)
May et al ²²	Validation	SHFM	n/a	n/a	n/a	n/a	Correlation (r=0.99)	†At 1 y: Total cohort=0.73 (0.71–0.75) Age >75 y (n=1339) =0.68 (0.65–0.72) LVEF >40% (n=1634)=0.66
Allen et al ²³	Validation	SHFM	n/a	n/a	n/a	n/a	Assessed graphically. Overestimated survival at 3 y by 8% (72% vs 80%).	At 1 y=0.73
Kalogeropoulos et al ²⁴ and Giamouzis et al ²⁵	s Validation	SHFM	n/a	n/a	n/a	n/a	H-L test, inadequate (P<0.05). Graphically, adequate after model recalibration	‡At 1 y: Total cohort (n=445)=0.78 ICD/CRT (n=316)=0.78 No ICD/CRT (n=129)=0.79 White (n=223)=0.78 Black (n=198)=0.79 (Continued

Table. Continued

Study	Derivation Validation	Model/Variables	Selection	Linear Gradient	Overfitting	Model Assumptions	Calibration	Discrimination (<i>c</i> -Statistic)
Levy et al ²⁶	Validation	SHFM and effect of IABP and inotropic support added from effect estimates obtained from previous studies		n/a	n/a	n/a		At 1 y=0.71
Gorodeski et al ¹⁸	Validation	SHFM	n/a	n/a	n/a	n/a	Tested graphically: overestimated survival in HT candidates and non-HT candidates	§At 1 y: In HT candidates=0.68 (0.63–0.74) In non-HT candidates=0.63 (0.57–0.69)
Goda et al ²¹	Validation	SHFM	n/a	n/a	n/a	n/a	n.r.	*At 1 y=0.73
Perrota et al ²⁷	Validation	SHFM	n/a	n/a	n/a	n/a	H-L test: <i>P</i> >0.2 at 1, 2, and 3 y	At 1 y=0.70 (0.61-0.79)
Haga et al28	Validation	SHFM	n/a	n/a	n/a	n/a	n.r.	Overall=0.68 (0.58-0.78)
Frankenstein et al ¹²	Derivation	 BNP 6'WT (different cutoff according to sex and β-blockers) 	Based on univariable analysis	n.r.	no	n.r.	n.r.	Overall: Unadjusted=0.76 Sex-adjusted=0.77 β-Blocker-adjusted=0.76 Sex-β-blocker-adjusted=0.77
	Validation	Frankenstein ¹²	n/a	n/a	n/a	n/a	n.r.	Unadjusted=0.66 Sex-adjusted=0.66 β-Blockers-adjusted=0.66 Sex-β-blockers-adjusted=0.68
Kramer et al ¹³	Derivation	PACE risk score Age >75 y LVEF <20% Creatinine PVD	Based on univariable analysis	n.r.	no	n.r.	n.r.	At 1 y=0.79
	Validation	PACE risk score	n/a	n/a	n/a	n/a	n.r.	At 1 y=0.69
Bilchick et al ¹⁴	Derivation	SHOCKED predictors Age NYHA LVEF COPD Diabetes mellitus Atrial fibrillation CKD	Based on clinical importance and statistical analysis	n.r.	no	n.r.	Correlation (<i>r</i> =0.89)	Overall=0.75 (0.75–0.76)
	Validation	SHOCKED predictors	n/a	n/a	n/a	n/a	Correlation (<i>r</i> =0.89) H-L test: <i>P</i> <0.001 at 2 and 3 y	Overall=0.74 (0.74-0.75)

6'WT indicates 6-minute walk test; ACEI, angiotensin-converting enzyme inhibitor; ARB, angiotensin II receptor blocker; BNP, brain natriuretic peptide; BP, blood pressure; CKD, chronic kidney disease; CMP, cardiomyopathy; COPD, chronic obstructive pulmonary disease; CRT, cardiac resynchronization therapy; ELITE2, Losartan Heart Failure Survival Study; HFSS, Heart Failure Survival Score; H-L, Hosmer–Lemeshow; HT, heart transplantation; IABP, intra-aortic balloon pump; ICD, internal cardiac defibrillator; IN-CHF, Italian Congestive Heart Failure Registry; IVCD, intraventricular conduction defect; LVEF, left ventricular ejection fraction; n/a, non applicable; n.r., not reported; NYHA, New York Heart Association; PVD, peripheral vascular disease; RCT, randomized controlled trial; RENAISSANCE, Randomized Etanercept North American Strategy to Study Antagonism of Cytokines; SHFM, Seattle Heart Failure Model; UW, University of Washington HF clinic; Val-HeFT, Valsartan Heart Failure Trial; and Vo₂, oxygen consumption.

*Goda et al²¹ reported that *c*-statistic was significantly higher (*c*-statistic=0.77 at 1 y) when HFSS and SHFM were used in a combined manner.

†Authors analyzed the additive discriminative value of creatinine, blood urea nitrogen (BUN), diabetes mellitus, and BNP (c-statistic=0.74, 0.74, 0.74, and 0.78, respectively).

‡Giamouzis et al²⁵ analyzed the additive of renal function and reported that renal function (BUN) did not significantly change discriminative capacity. §Authors analyzed the additive predicted value of BNP, BUN, and peak Vo, and reported nonsignificant improvement in *c*-statistic values.

However, this HFSS variant has not been further validated. Only 1 study¹8 assessed HFSS model calibration and reported that the model overestimated event-free survival by ≈20% in low-risk patients.

Seattle Heart Failure Model

The SHFM includes 10 continuous variables (age, LVEF, New York Heart Association class, systolic blood pressure, diuretic

dose adjusted by weight, lymphocyte count, hemoglobin, serum sodium, total cholesterol, and uric acid) and 10 categorical variables (sex, ischemic cardiomyopathy, QRS>120 ms, use of β -blockers, angiotensin-converting enzyme inhibitors, angiotensin receptor blockers, potassium-sparing diuretic, statins and allopurinol, and ICD/cardiac resynchronization therapy [CRT] status) in an equation that provides a continuous risk score for each patient, and which can be expressed as

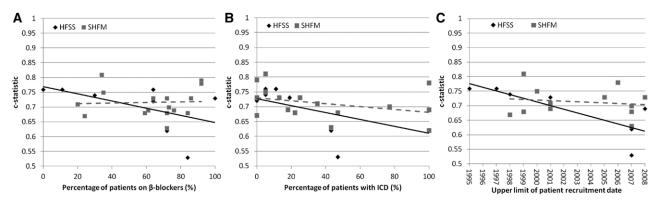


Figure 2. Model discrimination. Model discrimination according to the use of β-blockers (A), internal cardiac defibrillator (ICD; B), and study patients recruitment date (C). HFSS indicates Heart Failure Survival Score; and SHFM, Seattle Heart Failure Model.

predicted mean life expectancy or event-free survival at 1, 2, and 5 years. 11 This model was developed to predict a composite outcome of death, urgent heart transplantation, and ventricular assist device in 1125 patients with HF enrolled in the randomized controlled trial Prospective Randomized Amlodipine Survival Evaluation. The SHFM has been validated in 14 independent cohorts including 16057 patients with HF (4 cohorts including 8983 patients with HF were selected from randomized controlled trials [Table II in the online-only Data Supplement]). 11,18,22-28 The validation cohorts involve diverse populations with a mean age from 52 to 77 years, a higher proportion of males (61%-82%), and mean LVEF between 17% and 45%. In 4 cohorts, the used of β -blockers was 20% to 35%, and in the remaining cohorts was >60% (maximum of 92%). In 10 studies reporting ICD status, the use of ICD was <25% in 5 cohorts and >65% in 3 cohorts.

Model discrimination varied from poor to acceptable (0.63– 0.81), being at least modest (>0.70) in 7 (50%) cohorts of the 14 validation cohorts. There was a slight trend toward poorer discrimination in cohorts with higher use of ICD devices but was only weakly related to β-blocker use and recruitment date (Figure 2). Some studies^{18,22,25} have analyzed variations of the SHFM including other predictors, such as renal function, diabetes mellitus, peak oxygen consumption, and brain natriuretic peptide, and reported that discrimination did not improve significantly. However, May et al²² reported that discrimination was significantly improved from 0.72 to 0.78 when brain natriuretic peptide was added to the model. Model calibration was evaluated in most of the cohorts (Table) and showed a high correlation (r-coefficient >0.97) between observed and predicted survival. In 3 cohorts, calibration was assessed graphically by comparing observed and predicted event-free survival^{17,22,24}; the model overestimated event-free survival by ≈2% at 1 year and 10% at 5 years, more significantly in black and patients with ICD/CRT.²² The study by Kalogeropoulos et al24 reported inadequate model goodness-of-fit as assessed by the Hosmer-Lemeshow test.

Frankenstein et al's Model

This model includes 2 binary variables: brain natriuretic peptide and 6-minute walk test with different cutoffs depending on sex and use of β-blockers.¹² Patients can then be categorized into 3 groups (scores 0, 1, or 2). This model was derived from 636 patients with HF to predict all-cause mortality and

validated in an independent cohort of 676 patients with HF (mean age, 74 years; 76% male; 63% ischemic cardiomyopathy; 54% treated with β-blockers). Model discrimination in the validation cohort was poor, varying from 0.66 to 0.68 (Table). Model calibration was not reported.

PACE Risk Score

This model includes 4 binary variables: the presence of peripheral vascular disease, age >70 years, creatinine >2 mg/dL, and LVEF <20%, and it provides a continuous risk score for an individual patient from 0 to 5.13 This model was derived from 905 secondary and primary prevention patients with ICD to predict all-cause mortality and validated in an independent cohort of 1812 patients with ICD-HF (mean age, 64 years; 77% male; mean LVEF of 31%; and 58% had ischemic cardiomyopathy [Table II in the online-only Data Supplement]). Model discrimination in the validation cohort was poor with a c-statistic of 0.69 at 1 year (Table). Model calibration was not reported.

SHOCKED Predictors

This model includes 7 binary variables: age >75 years, New York Heart Association class >II, atrial fibrillation, chronic obstructive pulmonary disease, chronic kidney disease, LVEF <20%, and diabetes mellitus.¹⁴ This model provides a continuous risk score from 0 to 400 and estimates 1-, 2-, 3- and 4-year survival using a nomogram. This model was derived and validated from a cohort of Medicare beneficiaries receiving primary prevention ICD. The validation cohort included 27893 patients (39% of patients were >75 years, 75% male, 31% had LVEF <20%, and 63% had ischemic cardiomyopathy [Table II in the online-only Data Supplement]). Model discrimination in the validation cohort was modest with a c-statistic of 0.74 at 1 year (Table). Overall correlation between observed and predicted survival was high correlation (r-coefficient >0.89). However, model calibration, assessed by Hosmer-Lemeshow test, showed inadequate goodness-of-fit at 2 and 3 years.

Prediction Models Not Validated in an **Independent Cohort**

We identified 15 prediction models that were not validated in an external cohort. Tables IV, V, and VI in the online-only Data Supplement summarize the characteristics of studies included, the assessment of study quality, and model characteristics, respectively. These models include a wide variety of predictors tested in diverse HF populations. The number of predictors included ranged from 2 to 21. Seven models were derived from patients with reduced LVEF and 1 in patients with preserved LVEF. The remaining studies included patients with clinically diagnosed HF without considering a specific LVEF cutoff as an inclusion criterion. In 6 studies, internally validated by bootstrapping, model discrimination ranged from 0.74 to 0.85. The best discrimination (c-statistic, 0.85) was observed in the DSC (Dyssynchrony, posterolateral Scar location and Creatinine) index, a model derived from a selective cohort of patients with HF undergoing CRT implantation, which included some variables that are not routinely available: 1 binary variable, posterolateral scar location evaluated by cardiovascular magnetic resonance; and 2 continuous variables, tissue synchronization index measured by cardiovascular magnetic resonance and serum creatinine. The 5 studies that evaluated model calibration reported adequate performance.

Discussion

In this systematic review, we identified 20 event-free survival prediction models in ambulatory patients with HF. Only 25% (5 of 20 models) have been validated in external cohorts and only 2 models, the HFSS and the SHFM, have been validated in >2 independent cohorts, mostly reporting modest (0.70–0.80)-to-poor (<0.70) discrimination. Studies using the HFSS more frequently reported modest (>0.70) discrimination than cohorts evaluating the SHFM. However, HFSS performance showed a decline over time, whereas the SHFM had a relatively stable performance. Nonetheless, only 2 studies have directly compared models within the same population and reported that model discrimination was similar (c-statistic of 0.73 and 0.72 20 for the SHFM and 0.68 and 0.63 18 for the HFSS at 1 year).

Model discrimination represents the capacity of the model to differentiate patients who had the event from those who did not. The study by Goda et al²⁰ reported that discrimination was significantly higher (from 0.72–0.73 to 0.77 at 1 year) when HFSS and SHFM were used in a combined manner within the same model. May et al²² reported that the discrimination of the SHFM was significantly improved from 0.72 to 0.78 when brain natriuretic peptide was added to the model. As proposed by D'Agostino and Byung-Ho Nam,⁹ a model with discriminative capacity >0.70 has acceptable discrimination; a discriminative capacity >0.80 provides strong support to guide medical decision-making. Clearly, HFSS and SHFM have consistently demonstrated that their performance shows only modest discriminative capacity.

One potential reason for suboptimal performance is that the management and treatment of patients with HF has changed substantially in the past 2 decades. These models were derived from cohorts of patients recruited ≈ 20 years ago (1986–1991 for the HFSS and 1992–1994 for the SHFM).

As proposed by Moons et al,⁶ a good model should include variables that are believed to be associated with the outcome of interest. Koelling et al¹⁶ evaluated the association of the

7 predictors included in the HFSS model in patients treated with β-blockers and reported that only peak oxygen consumption and LVEF were factors independently associated with event-free survival. In addition, the directions of association of some predictors are opposite in the validation and derivation cohorts. For instance, the HFSS derivation study reported that the hazard ratio for 1 beat per minute increase in heart rate was 1.02 (95% confidence interval of 1.01–1.04), while in 2 validation cohorts 16,20 including a high proportion of patients treated with β-blockers (>70%), the hazard ratio was 0.98 (95% confidence interval, 0.97–1.01). This may partially explain the decline observed in the HFSS discriminatory capacity in more recent validation cohorts.

A similar situation is found with potassium-sparing diuretic use in the SHFM. Levy et al¹¹ imputed in the calculus of the score a hazard ratio of 0.74 for patients on potassium-sparing diuretics. Goda et al²⁰ reported a nonsignificant reverse effect of spironolactone in a contemporary cohort (hazard ratio, 1.20; 95% confidence interval, 0.86–1.48). Importantly, this tells us that predictors that were believed or found to be associated with mortality in patients with HF 20 years ago may not act similarly in contemporary patients with HF. This supports the need to develop and test an up-to-date prediction model.

Discrimination should not be reported in isolation because a poorly calibrated model can have the same discriminative capacity as a perfectly calibrated model.²⁹ One limitation of calibration is that assessment techniques do not allow for comparisons between models. In the validation cohorts, both the SHFM and the HFSS showed inadequate calibration attributable to the model overestimating survival in some groups of patients, including low-risk patients, blacks, and patients with ICD/CRT therapy.

Model ability to predict survival has not been compared with intuitive predictions of physicians. A study by Muntwyler et al³⁰ showed that primary care physicians overestimated mortality risk in patients with HF (1-year observed mortality of 13% versus physician estimate of 26%); this was more pronounced in stable New York Heart Association class II patients (1-year observed mortality of 6% versus physician estimated of 18%).

Whether these models may be used to guide or improve clinical practice remains underexplored. Vickers et al²⁹ have proposed the use of simple decision analytic techniques to compare prediction models in terms of their consequences. These techniques weight true and false-positive errors differently, to reflect the impact of decision consequences (ie, risks associated with heart transplantation or ventricular assist device versus risks associated with continuing medical therapy). Such decision analytic techniques may assist in determining whether clinical implementation of prediction models would do more good or more harm relative to current practice (physicians' predictions).

Should use and validation of these models continue? Or should we seek better models? There is no consensus on this issue among commentators. Researchers are pursuing both avenues, validating and supporting the use of the SHFM and HFSS as well as developing new models.

The performance of more recent models developed thus far, however, does not provide evidence that they will perform substantially better than older models. The 3 externally validated and recently published models¹²⁻¹⁴ have demonstrated poor-to-modest discrimination (between 0.66 and 0.74). Similarly, the 6 models that were validated by bootstrapping showed in general poor-to-modest discrimination. One of these 6 models provided high discriminatory capacity, but it was developed in a selected group of patients with HF undergoing CRT implantation and included 2 variables that are not easily measured (myocardial tissue synchronization index and scar location by cardiovascular magnetic resonance). The lack of external validation makes it difficult to assess how the performance of the model might be generalized to other populations, which clearly limits their clinical use. Discrimination estimated on a first sample is often higher than that on the subsequent samples.³¹

Other reasons potentially explaining the suboptimal performance of existing models may pertain to the presence of missing data and variable selection. For example, in cohorts validating the SHFM, the presence of missing data was as high as 100% for percentage of lymphocytes²⁶ or 65% for uric acid.²² Whether frequently missing or not easily available variables should be used to develop a score or should be incorporated to standard clinical practice will depend on the strength of the association between the predictors and outcome, the compromised model performance when the variables are not included in the final score and clinical resources. Nonetheless, adequate methods to deal with missing data, such as multiple imputation techniques, are important when evaluating model performance. The exclusion of cases because of missing information may lead to biased results.³²

Variable selection based on statistical significance may lead to suboptimal models. Other techniques, such as stability selection and subsampling, have demonstrated to yield more stable models based on a consistent selection of variables decreasing the chances of type I error.³³

As noticed in this review, the performance of predictive models has been traditionally evaluated by the c-statistic, which has been criticized as being insensitive in comparing models and for having limited direct clinical use. Reclassification tables, reclassification calibration statistic, and net reclassification and integrated discrimination improvements are recently developed methods to assess discrimination, calibration, and overall model accuracy. It has been shown that the use of these methods can better guide clinical decision-making by offering prognostic information at different risk strata. The use of these techniques is highly recommended during validation of existing or new models.

Conclusions

Optimal management of patients with HF requires accurate assessment of prognosis; however, making accurate assessment remains challenging. Among 5 externally validated prediction models, the HFSS and SHFM models demonstrated modest discriminative capacity and questionable calibration. The clinical impact of medical decision-making guided by the use of these models has not been explored. Given the limitation of current HF models, the development of a new model derived from contemporary patient cohorts is an appealing option. However, the development and reporting of new models should

be optimized by adhering to guidelines to guarantee model adequacy. In addition, new models should seek external validation of their generalizability and performance. Evaluation of the clinical impact of decisions based on models relative to current clinical practice would be enormously informative in determining their use in real-world clinical practice.

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Disclosures

None.

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CLINICAL PERSPECTIVE

Many models are available to predict adverse outcomes in patients with heart failure. Clinicians and researchers wishing to use prognostic models would benefit from knowledge of their characteristics and performance. Therefore, we performed a systematic review to identify studies evaluating risk prediction models for mortality in ambulatory patients with HF, to describe their performance and clinical applicability. This systematic review included 34 studies testing 20 models. Only 5 models were validated in an independent cohort: the Heart Failure Survival Score, the Seattle Heart Failure Model, the PACE risk score, a model by Frankenstein et al,¹² and the SHOCKED predictors. The Heart Failure Survival Score, validated in 8 cohorts, showed poor-to-modest discrimination (*c*-statistic, 0.56–0.79), being lower in the more recent validation studies possibly because of greater use of β-blockers and implantable cardiac defibrillators. The Seattle Heart Failure Model was validated in 14 cohorts describing poor-to-acceptable discrimination (0.63–0.81), remaining relatively stable over time. Both models reported adequate calibration, although overestimating survival in some specific populations. The other 3 models were validated in a cohort each, with poor-to-modest discrimination (0.66–0.74). There were no studies reporting the clinical impact of medical decision-making guided by the use of these models. In conclusion, externally validated HF models showed inconsistent performance. The Heart Failure Survival Score and Seattle Heart Failure Model demonstrated modest discrimination and questionable calibration. A new model derived from contemporary patient cohorts may be required for improved prognostic performance.

SUPPLEMENTAL METHODS

Appendix A: Literature Search Results

For: Ana Carolina Alba

Date Completed: 15 May 2012

The databases searched were:

- Ovid MEDLINE
- EMBASE
- CINAHL

RESULTS & STRATEGY USED:

Database: Ovid MEDLINE(R) <1946 to May Week 1 2012>

Search Strategy:

- 1 exp Heart Failure/ (76819)
- 2 ((heart or cardiac) adj2 failure).mp. (121311)
- 3 1 or 2 (121859)
- 4 predict:.mp. (756732)
- 5 validat:.tw. (180066)
- 6 scor:.tw. (404761)
- 7 observ:.mp. (2029286)
- 8 or/4-7 (3043863)
- 9 3 and 8 (28134)
- 10 exp Ambulatory Care/ (42583)
- 11 Outpatients/ (7351)
- 12 (ambulatory or stable or chronic or out-patient: or outpatient:).mp. [mp=title, abstract, original title, name of substance word, subject heading word, protocol supplementary concept, rare disease supplementary concept, unique identifier] (1246085)
- 13 10 or 11 or 12 (1246085)
- 14 9 and 13 (8814)
- (mortality or survival or death).mp. [mp=title, abstract, original title, name of substance word, subject heading word, protocol supplementary concept, rare disease supplementary concept, unique identifier] (1266793)
- 16 14 and 15 (3910)
- 17 statistics as topic/ or exp regression analysis/ (319979)

- 18 sn.fs. (425839)
- 19 statistic:.mp. (727873)
- 20 (logistic adj2 model:).mp. (85018)
- 21 (Likelihood adj2 function:).mp. (14814)
- 22 regression:.mp. (356421)
- 23 exp mathematical concepts/ (626843)
- 24 algorithm:.mp. (178754)
- 25 mathematic:.mp. (122305)
- 26 multivariate analysis/ (66832)
- 27 exp models, biological/ or exp models, statistical/ or logistic models/ (743997)
- 28 area under curve/ (21246)
- 29 or/17-28 (2456770)
- 30 "review"/ (1691446)
- 31 risk assessment/ or risk factors/ (590256)
- 32 evaluation.mp. (1000618)
- 33 exp Prognosis/ (930163)
- 34 prognostic factor:.mp. (47548)
- 35 8 or 31 or 32 or 33 or 34 (4702602)
- 36 3 and 13 and 15 and 35 (6181)
- 37 29 and 36 (2602)
- 38 30 and 36 (1361)
- 39 37 or 38 (3762)

Database: Embase <1974 to 2012 May 14>

Search Strategy:

- 1 exp heart failure/ (244924)
- 2 ((heart or cardiac) adj2 failure).mp. (207214)
- 3 1 or 2 (278699)
- 4 predict:.mp. (983853)
- 5 validat:.tw. (256546)
- 6 scor:.tw. (563146)
- 7 observ:.mp. (2609157)
- 8 risk assessment/ (285564)
- 9 risk factor/ (519981)
- 10 evaluation.mp. (1128376)
- 11 exp prognosis/ (388902)
- 12 prognostic factor:.mp. (67942)
- 13 or/4-12 (5511416)
- 14 3 and 13 (97265)
- 15 exp ambulatory care/ (35968)
- 16 outpatient/ (40332)
- 17 outpatient care/ (18777)
- 18 (ambulatory or stable or chronic or out-patient: or outpatient:).mp. (1647754)
- 19 15 or 16 or 17 or 18 (1647754)
- 20 14 and 19 (24318)
- 21 (mortality or survival or death).mp. (1806751)
- 22 20 and 21 (11345)
- 23 limit 22 to "review" (2010)
- 24 limit 23 to embase (1656)
- 25 exp statistics/ (272033)
- 26 exp regression analysis/ (179182)
- 27 statistic:.mp. (1196401)
- 28 (logistic adj2 model:).mp. (31580)
- 29 (Likelihood adj2 function:).mp. (782)
- 30 regression:.mp. (461195)
- 31 exp mathematical phenomena/ (2108262)
- 32 algorithm:.mp. (176636)
- 33 mathematic:.mp. (206662)
- 34 exp multivariate analysis/ (190591)
- 35 exp biological model/ (805064)

- 36 statistical model/ (88920)
- 37 area under the curve/ (55589)
- 38 or/25-37 (3631278)
- 39 22 and 38 (5358)
- 40 limit 39 to embase (4882)
- 41 24 or 40 (5993)

CINAHL Search Strategy Tuesday, May 15, 2012 1:44:33 PM

#	Query	Limiters/Expanders	Last Run Via	Results
S29	S18 or S28	Search modes - Boolean/Phrase	Interface - EBSCOhost Search Screen - Advanced Search Database - CINAHL	634
S28	S19 and S27	Search modes - Boolean/Phrase	Interface - EBSCOhost Search Screen - Advanced Search Database - CINAHL	569
S27	S20 or S21 or S22 or S23 or S24 or S25 or S26	Search modes - Boolean/Phrase	Interface - EBSCOhost Search Screen - Advanced Search Database - CINAHL	473798
S26	TX area under curve	Search modes - Boolean/Phrase	Interface - EBSCOhost Search Screen - Advanced Search Database - CINAHL	116
S25	(MH "Models, Theoretical+")	Search modes - Boolean/Phrase	Interface - EBSCOhost Search Screen - Advanced Search Database - CINAHL	52897
S24	(MH "Multivariate Analysis") OR (MH "Multivariate Analysis of Variance") OR (MH "Multivariate Analysis of Covariance")	Search modes - Boolean/Phrase	Interface - EBSCOhost Search Screen - Advanced Search Database - CINAHL	29451
S23	(MH "Mathematics+")	Search modes - Boolean/Phrase	Interface - EBSCOhost Search Screen - Advanced Search Database - CINAHL	291987
S22	TX statistic* or TX logistic N2 model* or TX likelihood N2 function* or TX regression or TX algorithm* or TX mathematic*	Search modes - Boolean/Phrase	Interface - EBSCOhost Search Screen -	428036

			Advanced Search Database - CINAHL	
S21	(MH "Regression+")	Search modes - Boolean/Phrase	Interface - EBSCOhost Search Screen - Advanced Search Database - CINAHL	109567
S20	(MH "Statistics+")	Search modes - Boolean/Phrase	Interface - EBSCOhost Search Screen - Advanced Search Database - CINAHL	282038
S19	S16 and S17	Search modes - Boolean/Phrase	Interface - EBSCOhost Search Screen - Advanced Search Database - CINAHL	1136
S18	S16 and S17	Limiters - Publication Type: Review Search modes - Boolean/Phrase	Interface - EBSCOhost Search Screen - Advanced Search Database - CINAHL	73
S17	TX mortality or TX survival or TX death	Search modes - Boolean/Phrase	Interface - EBSCOhost Search Screen - Advanced Search Database - CINAHL	158882
S16	S11 and S15	Search modes - Boolean/Phrase	Interface - EBSCOhost Search Screen - Advanced Search Database - CINAHL	2698
S15	S12 or S13 or S14	Search modes - Boolean/Phrase	Interface - EBSCOhost Search Screen - Advanced Search Database - CINAHL	175366
S14	TX ambulatory or TX stable or TX chronic or TX out-patient* or TX outpatient*	Search modes - Boolean/Phrase	Interface - EBSCOhost Search Screen - Advanced Search Database - CINAHL	171927

S13	(MH "Outpatients") OR (MH "Outpatient Service")	Search modes - Boolean/Phrase	Interface - EBSCOhost Search Screen - Advanced Search Database - CINAHL	29357
S12	(MH "Ambulatory Care") OR (MH "Ambulatory Care Facilities+") OR (MH "Ambulatory Care Nursing")	Search modes - Boolean/Phrase	Interface - EBSCOhost Search Screen - Advanced Search Database - CINAHL	13447
S11	S9 and S10	Search modes - Boolean/Phrase	Interface - EBSCOhost Search Screen - Advanced Search Database - CINAHL	8549
S10	S3 or S4 or S5 or S6 or S7 or S8	Search modes - Boolean/Phrase	Interface - EBSCOhost Search Screen - Advanced Search Database - CINAHL	602415
S9	S1 or S2	Search modes - Boolean/Phrase	Interface - EBSCOhost Search Screen - Advanced Search Database - CINAHL	20275
S8	TX "prognostic factor*"	Search modes - Boolean/Phrase	Interface - EBSCOhost Search Screen - Advanced Search Database - CINAHL	2789
S7	(MH "Prognosis+")	Search modes - Boolean/Phrase	Interface - EBSCOhost Search Screen - Advanced Search Database - CINAHL	119023
S6	TX evaluation	Search modes - Boolean/Phrase	Interface - EBSCOhost Search Screen - Advanced Search Database - CINAHL	263029
S5	(MH "Risk Factors+")	Search modes - Boolean/Phrase	Interface - EBSCOhost Search Screen -	62487

			Advanced Search Database - CINAHL	
S4	(MH "Risk Assessment")	Search modes - Boolean/Phrase	Interface - EBSCOhost Search Screen - Advanced Search Database - CINAHL	27594
S3	TX predict* or TX validat* or TX scor* or TX observ*	Search modes - Boolean/Phrase	Interface - EBSCOhost Search Screen - Advanced Search Database - CINAHL	276104
S2	TX heart N2 failure or TX cardiac N2 failure	Search modes - Boolean/Phrase	Interface - EBSCOhost Search Screen - Advanced Search Database - CINAHL	20263
S1	(MH "Heart Failure+")	Search modes - Boolean/Phrase	Interface - EBSCOhost Search Screen - Advanced Search Database - CINAHL	

Appendix B. Study eligibility form¹

Reviewer:	XX	ZZ	NN	
				•
Article ID:				
Reference #:	Auth	or: Jo	ournal:	Year:

Popula	Population ² :						
•	Ambulatory heart failure patients	YES	NO				
•	Adults (≥ 19 years old)	YES	NO				

Predictive model ³ :					
 ≥ 2 predictors or Validation study of pre-existing score 	YES	NO			
Report of score formula or coefficients and intercept	YES	NO			
 Assessment of discrimination and/or calibration 	YES	NO			

I	Outcomes reported:							
	•	Mortality or composite outcome including mortality	YES	NO				
I	•	>30 deaths	YES	NO				

Study	Study design:						
•	Cohort study (prospective or retrospective) or						
•	Randomized control trial or	YES	NO				
•	Meta-analysis						

Duplic	Duplicated population:										
•	If duplicated, does this study report new information on model	YES	NO								
	performance?										

Study inclusion:	
All the answers are YES	INCLUDE
Any answer is NO	EXCLUDE

¹ If any response to the above questions is unclear, mark YES.
² If a study included hospitalized patients or transplant or VAD patients, consider as NO.

³ Any type of predictor, including but not limited to clinical characteristics, laboratory values, test results and any other clinical event, such as hospital admissions, ICD shocks, etcetera.

SUPPLEMENTAL TABLES

Supplemental Table 1. Aspects considered in the assessment of model adequacy and performance

Item	Description
Selection of the	A good model should clearly state how predictors were selected. Potential candidate
predictors	predictors may be chosen according to correlation with the outcome of interest
	explored in univariable analysis or based on previous knowledge. Whether one
	approach is better than the other is a matter of unresolved discussion. The former
	may include predictors that are not necessarily casual while the latter requires robust
	knowledge on the field of study.
Coding of the	The proper reporting of the coding of variables is important because the effect of an
predictors	independent variable on the outcome variable depends on the corresponding units of
	measurement and the manner in which the variable was coded. Articles were
	considered to properly report the coding of variables if the method of coding for all of
	the variables that remained in the final statistical model could easily be determined
	or were referenced anywhere in the article.
Nonconformity	If the manuscript did not report determining the impact of each explanatory variable
to a Linear	separately in zones of ranked data or mentioned that conformity to a linear gradient
Gradient	was addressed, this item was coded as not reported.
Over-fitting	Risk estimates may be unreliable if the multivariable model includes too many
	independent variables and too few outcome events, they may represent spurious
	associations or the effects may be estimated with low precision. According to Peduzzi
	et al [1], we categorized the articles with a ratio of < 10:1 (10 outcome events for
	each single explanatory variable in the final model) as an over-fitted.

Analysis of	Violation of model assumptions, such as the proportional hazards assumption in the
statistical model	case of Cox method, may lead to unreliable effect estimates. If a manuscript did not
assumption	state exploring model assumptions and that they were held in the final proposed
	model, this item was coded as not reporting model assumptions.
Discrimination	Discrimination expresses to what extent the model is capable of differentiating
	patients who had the event from those who did not. It is commonly assessed using
	the c-statistic test, which is equivalent to the area under the receiver operating
	characteristic (ROC) curve [2]. The ROC curve is a plot of sensitivity versus 1-
	specificity, which are calculated for each value of the predicted risk as a possible cut-
	off value. A c-statistic of 0.50 indicates that the model performs no better than
	chance; a c-statistic of 0.50 to 0.70 indicates poor discrimination; a c-statistic of 0.70
	to 0.80 indicates modest discriminative ability; and a c-statistic of greater than 0.80
	indicates aceptable discriminative ability [2].
Calibration or	The calibration or goodness of fit of a model measures how well the model describes
goodness of fit	the response variable. Goodness-of-fit involves investigating how close values
	predicted by the model are to the observed values. It can be assessed using different
	methods (i.e., Hosmer-Lemeshow test or deviance, Cox-Snell analysis, correlation
	between observed vs. predicted events).

References of Supplemental Table 1:

- 1. Peduzzi P, Concato J, Feinsten AR, Holford TR. Importance of events per independent variable in proportional hazards regression analysis II. Accuracy and precision of regression estimates. J Clin Epidemiol 1995;48:1503-10.
- 2. D'Agostino RB, Byung-Ho Nam. Evaluation of the performance of survival analysis models: Discrimination and calibration measures. In: Handbook of Statistics v23: Advances in survival analysis, by Balakrishnan N, Rao CR. 2004.

Supplemental Table 2. Characteristics of the population of studies included

Study	Model's	Derivation/		Population									Events	
	name	Validation	Source	Inclusion	Time	N	Mean	%	Mean	%	% β-	%	Definition	n
		study		criteria	frame		Age	male	LVEF	ischemic	blocker	ICD		
Aaronson [1]	HFSS	Derivation	Single	LVEF <40%	1986-	268	50	80	20	45	10	n.r.	Death and	109
1997			center	Age <70 years	1991								urgent HTx	
USA		Validation	Single		1993-	199	52	81	22	47	11	n.r.		~60
			centre		1995									
Zugck [2]	HFSS	Validation	Single	NYHA I-III	1995-	208	54	82	22	29	30	n.r.	Death	52
2001			center	LVEF <40%	1998									
Germany				Age <70 years					***************************************					
Koelling [3]	HFSS	Validation	Single	LVEF <40%	1994-	320	52	74	23	52	10	11	Death,	64
2004			center	CP study	1997								urgent HTx	
USA					1999-	187	54	76	21	56	72	19	and VAD	30
					2001									
Parikh [4]	HFSS	Validation	Single	HF	n.r.	396	70	75	30	50	64	n.r.	Death,	111
2009			center	Age >65 years									urgent HTx	
USA				CP study									and VAD	
Gorodeski [5]	SHFM	Validation	Single	Referred for	2004-	215	55	77	20	55	80	78	Death,	157
2010	HFSS		centre	HTx assessment	2007								urgent HTx	
USA				THE DESIGNATION OF THE PARTY OF									and VAD	

Supplemental Table 2. Continued

Study	Model's	Derivation/				Po	pulation						Events	
	name	Validation	Source	Inclusion	Time	N	Mean	%	Mean	%	% β-	%	Definition	n
		study		criteria	frame		Age	male	LVEF	ischemic	blocker	ICD		
Goda [6-8]	HFSS	Validation	Single	Referred for	1993-	715	54	65	22	40	71	49	Death,	354
2010	SHFM		center	HTx assessment	2008								urgent HTx	
USA	3 papers			THE THEORET STATES AND THE STATES AN									and VAD	
Levy [9]	SHFM	Derivation	PRAISE-1	LVEF <30%	1992-	1125	65	76	21	64	0	0		403
2006			Trial		1994									
USA			ELITE2	LVEF <40%	1997-	2987	71	69	31	74	24	0		505
			Trial	Age >60 years	1998									
			RENAISSA	LVEF <30%	1999-	925	62	78	22	61	61	18	Darth	179
			NCE trial	NYHA II-IV	2001								Death, urgent HTx	
		Validation [#]	Val-HeFT	LVEF <40%	1997-	5010	63	80	27	58	34	n.r.		979
		validation	Trial	NYHA II-IV	1999								and VAD	
			IN-CHF	HF patients	1995-	872	64	76	35	47	35	n.r.		115
			Registry		n.r.									
			UW	HF patients	n.r.	148	53	78	27	34	72	22		48
			Cohort											

Supplemental Table 2. Continued

Study	Model's	Derivation/				Po	pulation						Events	;
	name	Validation	Source	Inclusion	Time	N	Mean	%	Mean	%	% β-	%	Definition	n
		study		criteria	frame		Age	male	LVEF	ischemic	blocker	ICD		
May [10]	SHFM	Validation	Single	Hospitalized HF	1993-	4077	67	61	45	60	77	13	Death,	2142
2007			centre	patients	2005								urgent HTx	
USA													and VAD	
Allen [11]	SHFM	Validation	Single	HF patients	2004-	122	61	62	26	38	86	25	Death	35
2008			centre		2008									
USA														
Kalogeropoulos	SHFM	Validation	Single	LVEF <30%	2000-	445	52	69	18	38	92	68	Death,	109
[12] Giamouzis			centre	NYHA II-IV	2006								urgent HTx	
[13] 2009 USA													and VAD	
Levy [14]	SHFM	Validation	REMATCH	HF non-HTx	1998-	61	68	82	17	69	20	35	Death	56
2009			trial	candidates	2001									
Atlanta, USA				(medical										
				treatment arm)										
Perrota [15]	SHFM	Validation	Single	NYHA I-III	2000-	342	71	79	26	52	73	77	Death and	86
2012			centre	LVEF <35%	2007								urgent HTx	
Italy				CRT implant										

Supplemental Table 2. Continued

Study	Model's	Derivation/				Ро	pulation						Events	
	name	Validation	Source	Inclusion	Time	N	Mean	%	Mean	%	% β-	%	Definition	n
		study		criteria	frame		Age	male	LVEF	ischemic	blocker	ICD		
Haga [16]	SHFM	Validation	Single	NYHA III-IV	n.r.	138	77	66	n.r.	68	59	n.r	Death	43
2012			centre	No HF										
UK				admissions for										
				6 weeks										
Frankenstein	-	Derivation	Single	LVEF <40%	1995-	636	56	81	28	32	78	n.r	Death	151
[17]			center		2005									
2011		Validation			2001-	676	74	76	34	63	54	n.r.		160
Germany					2005									
Kramer [18]	PACE risk	Derivation	Multi-	Primary and	2001-	905	65	78	31	59	n.r.	100	Death	125
2012	score		center	secondary	2008									
USA		Validation		prevention	2001-	1812	64	77	31	58	n.r.	100		296
				ICD patients	2008									
Bilchick [19]	SHOCKED	Derivation	Multi-	Primary	2005-	17991	n.r.	77	n.r.	59	79	100	Death	6741
2012	predictors		center	prevention	2006									
USA		Validation	(Medicare	ICD patients	2005-	27893	n.r.	75	n.r.	63	n.r.	100		8595
			database)		2007									

HFSS, Heart Failure Survival Score; LVEF, left ventricular ejection fraction; HTx, heart transplantation; NYHA, New York Heart Association; CP, cardio-pulmonary; VAD, ventricular assist device; SHFM, Seattle Heart Failure Model; MI; myocardial infarction; PRAISE, Prospective Randomized Amlodipine Survival Evaluation; ELITE2, Losartan Heart Failure Survival Study; RENAISSANCE, Randomized Etanercept North American Strategy to Study Antagonism of Cytokines; IN-CHF, Italian Congestive Heart Failure Registry; UW, University of Washington HF clinic; CRT, cardiac resynchronization therapy; HF, heart failure; ICD, internal cardiac defibrillator; n.r., not reported.

Supplemental Table 3. Assessment of study quality

Study	Derivation	Model	Patient	Data collection	Missing data	Loss of
	Validation		selection			follow up
Aaronson 1997 [1]	Derivation	HFSS	n.r.	Retrospective	n.r.	1-3%
	Validation	HFSS	n.r.	Retrospective	n.r.	1-3%
Zugck 2001 [2]	Validation	HFSS	n.r.	Retrospective	n.r.	0%
Koelling 2004 [3]	Validation	HFSS	n.r.	Retrospective	0%	0%
Parikh 2009 [4]	Validation	HFSS	n.r.	Retrospective	36% of patients excluded	0%
Gorodeski 2010 [5]	Validation	HFSS	Consecutive	Retrospective	Peak VO ₂ = 36%. Imputed by multiple	n.r.
					imputation	
Goda 2010 [6] and	Validation	HFSS	Consecutive	Retrospective	18 patients excluded	0%
2011 [7,8]						
Levy 2006 [9]	Derivation	SHFM	RCT	Prospective	n.r.	n.r.
	PRAISE-1					
	Validation	SHFM	RCT	Prospective	n.r.	n.r.
	ELITE2					
	Validation	SHFM	RCT	Prospective	n.r.	n.r.
	Val-HeFT					

Supplemental Table 3. Continued.

Study	Derivation	Model	Patient	Data collection	Missing data	Loss of
	Validation		selection			follow up
Levy 2006 [9]	Validation	SHFM	n.r.	Prospective	n.r.	n.r.
	UW					
	Validation	SHFM	RCT	Prospective	n.r.	n.r.
	RENAISSANCE					
	Validation	SHFM	Registry	Prospective	n.r.	n.r.
	IN-CHF					
May 2007 [10]	Validation	SHFM	Consecutive	Prospective	NYHA=72%	0%
					Lymphocytes=35%	
					Uric acid=66%	
					LVEF=25%	
					Cholesterol=20%	
					Imputed using multiple regression	
Allen 2008 [11]	Validation	SHFM	Consecutive	Prospective	Imputed with the mean	0%

Supplemental Table 3. Continued

Study	Derivation	Model	Patient	Data collection	Missing data	Loss of
	Validation		selection			follow up
Kalogeoropoulos [12]	Validation	SHFM	Consecutive	Retrospective	Exclusion of patients with >2 missing	0%
and Giamouzis [13]					variables. The rest were imputed with	
2009					the mean (lymphocytes=71%).	
Levy 2009 [14]	Validation	SHFM	RCT	Prospective	Lymphocytes imputed by multiple	0%
					regression. Uric acid, cholesterol and	
					diuretic dose were imputed from a	
					comparable group of patients from	
					SHFM cohort.	
Gorodeski 2010 [5]	Validation	SHFM	Consecutive	Retrospective	Uric acid = 64%	n.r.
					Cholesterol = 11%	
					Lymphocytes = 10%	
					Imputed by multiple imputation	
Goda 2011 [8]	Validation	SHFM	Consecutive	Retrospective	In 38% patients, imputed with the	0%
					mean	
Perrota 2012 [15]	Validation	SHFM	n.r.	Retrospective	Imputed with the mean	n.r.

Supplemental Table 3. Continued.

Derivation	Model	Patient	Data collection	Missing data	Loss of
Validation		selection			follow up
Validation	SHFM	n.r.	Retrospective	n.r.	n.r.
Derivation	-	Consecutive	Retrospective	n.r.	n.r.
Validation		Consecutive	Retrospective	n.r.	n.r.
Derivation	PACE risk	Consecutive	Retrospective	n.r.	n.r.
	score				
Validation		Consecutive	Retrospective	n.r.	n.r.
Derivation	SHOCKED	Consecutive	Prospective	n.r.	n.r.
	predictors				
Validation		Consecutive	Prospective	n.r.	n.r.
	Validation Validation Derivation Validation Derivation Validation Derivation	Validation Validation Derivation Derivation PACE risk score Validation Derivation SHOCKED predictors	ValidationselectionValidationSHFMn.r.Derivation-ConsecutiveValidationConsecutiveDerivationPACE risk scoreConsecutiveValidationConsecutiveDerivationSHOCKED predictorsConsecutive	ValidationselectionValidationSHFMn.r.RetrospectiveDerivation-ConsecutiveRetrospectiveValidationConsecutiveRetrospectiveDerivationPACE risk scoreConsecutiveRetrospectiveValidationConsecutiveRetrospectiveDerivationSHOCKED predictorsConsecutiveProspective	Validation selection Validation SHFM n.r. Retrospective n.r. Derivation - Consecutive Retrospective n.r. Validation Consecutive Retrospective n.r. Derivation PACE risk score Consecutive Retrospective n.r. Validation Consecutive Retrospective n.r. Derivation SHOCKED consecutive Prospective n.r.

HFSS, Heart Failure Survival Score; peak VO₂, peak oxygen consumption; RCT, randomized controlled trial; SHFM, Seattle Heart Failure Model; PRAISE, Prospective Randomized Amlodipine Survival Evaluation; ELITE2, Losartan Heart Failure Survival Study; RENAISSANCE, Randomized Etanercept North American Strategy to Study Antagonism of Cytokines; IN-CHF, Italian Congestive Heart Failure Registry; UW, University of Washington HF clinic; LVEF, left ventricular ejection fraction; n.r., not reported.

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Supplemental Table 4. Characteristics of the population of studies included

Study	Model	Derivation/	Population										Events	
	name	Validation	Source	Inclusion	Time	N	Mean	%	Mean	%	% β-	%	Definition	n
		study		criteria	frame		Age	male	LVEF	ischemic	blocker	ICD		
Kearney	-	Derivation	Heart	Clinically	1993-	553	63	76	42	79	8	n.r.	Death	201
2003 [1]			study	diagnosed	1995									
UK				HF NYHA I-III										
Rickli 2003 [2]	-	Derivation	Single	LVEF<40%	n.r.	202	52	86	28	53	45	n.r	Death and	59
Switzerland			center	CP study									urgent HTx	
Adlam	-	Derivation	Single	Clinically	1995-	532	75	41	45	41	14	n.r.	Death	190
2005 [3]			centre	diagnosed	1998									
UK				HF										
Pocock 2006	CHARM	Derivation	CHARM	Clinically	1999-	7599	65	68	39	57	n.r.	n.r.	Death	1831
[4] UK			trial	diagnosed	2003									
				HF										
Myers	CPX score	Derivation	Multi-	Clinically	1993-	710	56	80	34	39	63	n.r.	Death,	110
2008 [5]			center	diagnosed	2007								urgent HTx	
Italy				HF									and VAD *	

Supplemental Table 4. Continued.

Study	Model	el Derivation/	Population										Events	
	name	Validation	Source	Inclusion	Time	N	Mean	%	Mean	%	% β-	%	Definition	n
		study		criteria	frame		Age	male	LVEF	ischemic	blocker	ICD		
Huynh	-	Derivation	Single	HF patients	1990-	282	80	34	42	54	n.r.	n.r.	Death	43
2008 [6]			center	Age >70 years	1994									
USA														
Wedel	CORONA	Derivation	CORONA	LVEF <40%	2003-	3342	72	73	32	100	78	2.3	Death *	934
2009 [7]	score		trial	NYHA II-IV	2005									
Europe														
Leyva	DSC index	Derivation	Single	LVEF<35%	2001-	148	68	77	23	62	55	0	CV Death	37
2009 [8]			center	NYHA III-IV	2008									
UK				CRT implant										
Vazquez	MUSIC	Derivation	Multi-	Clinically	2003-	992	65	72	37	46	68	n.r.	Death *	267
2009 [9]	score		centre	diagnosed HF	2004									
Spain				NYHA II-IV										
Komajda	-	Derivation	I-	LVEF >45%	2003-	4128	72	40	59	25	n.r.	n.r.	Death *	881
2011 [10]			PRESERVE	NYHA II-IV	2007									
France			trail	Age >50 years										

Supplemental Table 4. Continued

Study	Model's	Derivation/	Population										Events	
	name	Validation	Source	e Inclusion criteria	Time frame	N	Mean Age	% male	Mean LVEF	% ischemic	% β-	% ICD	Definiti on	N
		study									blocker			
Subramanian	VEST score	Derivation	VEST trail	LVEF <30%	1995-	963	62	78	21	57	n.r.	n.r.	Death *	172
2011 [11]				NYHA III-IV	1996									
USA				manus										
O'Connor	HF-ACTION	Derivation	HF-	LVEF <35%	2003-	2331	59	72	25	54	95	40	Death *	387
2012 [12]	score		ACTION	NYHA II-IV	2007									
USA			trail											
Herrmann		Derivation	Single	LVEF <40%	n.r.	114	63	n.r.	29	n.r.	4	n.r.	Death	31
2012 [13]			centre	HF										
UK				symptoms										
Scrutinio		Derivation	Single	LVEF <40%	2001-	802	64	79	28	50	73	n.r.	Death	301
2012 [14]			centre	HF	2007									
Italy				symptoms										
Pocock		Derivation	Multi-	Clinically	n.r.	39372	67	67	35	53	34	n.r.	Death	15851
2012 [15]			centre	diagnosed										
Europe				HF										

HF, heart failure; NYHA, New York Heart Association; CP, cardio-pulmonary; LVEF, left ventricular ejection fraction; HTx, heart transplantation; VAD, ventricular assist device; CV, cardiovascular; n.r., not reported.

Supplemental Table 5. Assessment of study quality

Study	Derivation	Model	Patient	Data collection	Missing data	Loss of
	Validation		selection			follow up
Kearney 2003 [1]	Derivation		n.r.	Prospective	Multiple regression	n.r.
Rickli 2003 [2]	Derivation		Consecutive		n.r.	n.r.
Adlam 2005 [3]	Derivation		Consecutive	Prospective	Excluded	0%
Pocock 2006 [4]	Derivation	CHARM	RCT cohort	Prospective	n.r.	n.r.
Myers 2008 [5]	Derivation	CPX score	n.r.	Prospective	n.r.	n.r.
Huynh 2008 [6]	Derivation		RCT cohort	Prospective	n.r.	n.r.
Wedel 2009 [7]	Derivation	CORONA	RCT cohort	Prospective	Excluded	n.r.
Leyva 2009 [8]	Derivation	DSC index	Consecutive	Prospective	0%	0%
Vazquez 2009 [9]	Derivation	MUSIC score	Consecutive	Prospective	Imputed with the mean	1.1%
Komajda 2011 [10]	Derivation		RCT cohort	Prospective	Excluded	n.r.
Subramanian 2011 [11]	Derivation	VEST	RCT cohort	Prospective	19% of patients excluded	n.r.

Study	Derivation	Model	Patient	Data collection	Missing data	Loss of
	Validation		selection			follow up
O'Connor 2012 [12]	Derivation	HF-ACTION	RCT cohort	Prospective	Hemoglobin= 24%	n.r.
					Urea= 13%	
					Sodium= 11%	
					Creatinine= 10%	
					MR= 8%	
					Multiple imputation	
Herrmann 2012 [13]	Derivation		n.r.	Prospective	n.r.	n.r.
Scrutinio 2012 [14]	Derivation		Consecutive	Prospective	0%	0%
Pocock 2012 [15]	Derivation		Meta-analysis	Prospective and	Multiple imputation	0%
			on RCT and	retrospective		
			observational			
			studies			

LVEF, left ventricular ejection fraction; ICD, internal cardiac defibrillator; HFSS, Heart Failure Survival Score; HTx, heart transplantation; VAD, ventricular assist device; NYHA, New York Heart Association; MFH; metabolic, functional, hemodynamic; CPX, cardiopulmonary exercise test; MRT, mean response time; SHFM, Seattle Heart Failure Model; MI; myocardial infarction; DSC, Dyssynchrony, posterolateral Scar location and Creatinine; CRT, cardiac resynchronization therapy; CV, cardiovascular; n.r., not reported.

Supplemental Table 6. Model derivation and performance

Study	Derivation	Model/ Variables	Selection	Linear	Over-	Model	Calibration	Discrimination
	Validation			Gradient	fitting	assumptions		(c-statistic)
Zugck 2001 [15]	Derivation	• LVEF	n.r.	n.r.	No	n.r.	n.r.	Overall = 0.84 (0.80-0.88) or
		• Peak VO ₂ or 6'WT						0.83 (0.79-0.87)
Kearney	Derivation	• Sodium	Based on	n.r.	Yes (201	Held	n.r.	* Binary predictors= 0.74
2003 [1]		Creatinine	univariable		events			(0.70-0.78)
		• CT ratio	analysis		and 30			Continuous predictors=
		• QRS dispersion			variables			0.78 (0.74-0.82)
		• QT			tested)			
		Non-sustained VT						
		• LVH by ECG						
		• SDNN						
	Validation by	Kearney	n/a	n/a	n/a	n/a	n.r	n.r.
	bootstrap	2003						

Study	Derivation	Model/ Variables	Selection	Linear	Over-	Model	Calibration	Discrimination
	Validation			Gradient	fitting	assumptions		(c-statistic)
Rickli 2003 [2]	Derivation	• Predicted peak	Based on	n.r.	No	n.r.	n.r.	At 1 year=0.86 (0.82-0.90)
		VO ₂	univariable					
		• MRT >50 seconds	analysis					
		• Systolic BP						
Adlam 2005 [3]	Derivation	• BNP	Based on	n.r.	No	Held	n.r.	Overall = 0.76
		• Age	univariable					
		• Sex	analysis					
		• Diabetes	using					
		• CVA	bootstrap					
		Abnormal ECG	estimated					
	Validation by	Adlam	n/a	n/a	n/a	n/a	n.r.	Overall = 0.75
	bootstrap	2005						

Study	Derivation	Model/ Variables	Selection	Linear	Over-	Model	Calibration	Discrimination
	Validation			Gradient	fitting	assumptions		(c-statistic)
Pocock 2006 [4]	Derivation	CHARM:	Probably	n.r.	No	n.r.	Graphically	At 2 years = 0.75
		• Age	on clinical				observed vs.	In preserved EF = 0.74
		• Sex	importanc				predicted	In low-EF=0.76
		• Diabetes	e. Forward				survival by	
		• LVEF	selection				deciles.	
		• NYHA					Under-	
		Cardiomegalia					estimated	
		• Time HF diagnose					survival at 3	
		• Prior HF					years	
		admission						
		• BMI						
		• Diastolic BP						
		• Smoking						
		• BBB						
		• Previous MI						

bootstrap							
Validation by	CHARM	n/a	n/a	n/a	n/a	n.r.	At 2 years = 0.75
	Candesartan						
	• Rest dyspnea						
	 Atrial fibrillation 						
	regurgitation						
	• Mitral						
	• Heart Rate						
	edema						
	• Pulmonary						
	• Edema						
	crackles						
	Pulmonary						

Study	Derivation	Model/ Variables	Selection	Linear	Over-	Model	Calibration	Discrimination
	Validation			Gradient	fitting	assumptions		(c-statistic)
Myers 2008 [5]	Derivation	CPX score:	Not clearly	n.r.	No	Held	n.r.	n.r.
		• OUES>1.4	stated					
		• VE/VCO2 >34						
		• peak VO2<14						
		• HR recovery <6						
		beats at 1minute						
		• PetCO2						
		<33mmHg						
	Validation by	CPX score	n/a	n/a	n/a	n/a	n.r.	‡ Overall = 0.77
	bootstrap							
Huynh 2008 [6]	Derivation	• Urea	Based on	n.r.	Yes	n.r.	n.r.	At 6 months=0.80
		Systolic BP	univariable		(43			
		• PVD	analysis.		events			
		• Sodium			and 15			
					variables)			

Study	Derivation	Model/ Variables	Selection	Linear	Over-	Model	Calibration	Discrimination
	Validation			Gradient	fitting	assumptions		(c-statistic)
	Validation by	Huynh	n/a	n/a	n/a	n/a	n.r.	n.r.
	bootstrap	2008						
Wedel 2009 [7]	Derivation	CORONA:	Not clearly	n.r.	No	n.r.	n.r.	Overall mortality=0.72
		• BNP	stated					HF mortality=0.80
		• Age						
		• Diabetes						
		• LVEF						
		• BMI						
		• Sex						
		• CABG						
		Atrial fibrillation						
		• NHYA						
		• Apo-A1						
		Creatinine						
		• PVD						

		Heart rate						
		• MI						
Leyva 2009 [8]	Derivation	DSC index:	Based on	Checked	No	Held	Correlation	At 1 year = 0.88
		Dyssynchrony	previous	by			(r=0.93)	At 1 year = 0.87
		Scar location	reports	martingal				
		Creatinine		e residuals				
	Validation by	DSC index	n/a	n/a	n/a	n/a	***	Overall=0.85
	bootstrap							
Vazquez 2009	Derivation	MUSIC score:	Based on	n.r.	No	n.r	Correlation	Overall mortality=0.76
[9]		• Prior MI, stroke	previous				(r=0.99)	Cardiac mortality=0.78
		or limb ischemia	knowledge					HF mortality=0.80
		• Left atrium	and <5%					Sudden death=0.77
		size>26mm/m2	missing					
		• LVEF<35%	data					
		• LBBB or IVCD						
		(QRS>110)						
		• non-sustained VT						
		or frequent						

		extra-beats						
		• GFR <60ml/min						
		• BNP>1000pg/dl						
		• Troponin posit						
		• Sodium						
		<138meq/L						
	Validation by	MUSIC score	n/a	n/a	n/a	n/a	n.r.	Overall mortality=0.77
	bootstrap							Cardiac mortality=0.78
								HF mortality=0.80
								Sudden death=0.78
1			1	1	1	1	I	i e

Study	Derivation	Model/ Variables	Selection	Linear	Over-	Model	Calibration	Discrimination
	Validation			Gradient	fitting	assumptions		(c-statistic)
Komajda 2011	Derivation	• BNP	Based on	n.r.	No	n.r.	Graphically	Overall=0.74
[10]		• Age	univariable				observed vs.	
		• Diabetes	analysis				predicted =	
		• LVEF					Adequate	
		Heart rate						
		• Previous hospital						
		admission						
		• Quality of life						
		• COPD or asthma						
		• Ischemic CMP						
		• MI						
	Validation by	Kornajda 2011	n/a	n/a	n/a	n/a	n.r.	Overall=0.74
	bootstrap							

Study	Derivation	Model/ Variables	Selection	Linear	Over-	Model	Calibration	Discrimination
	Validation			Gradient	fitting	assumptions		(c-statistic)
Subramanian	Derivation	VEST:	Based on	n.r.	Yes	n.r.	n.r.	Overall=
2011 [11]		Model:1	univariable		(172			Model 1: 0.73
		• BUN	analysis		events			Model 2: 0.74
		• LVEF			and 19			Model 3: 0.81
		• Lymphocytes			variables			
		• CT radio			tested)			
		Model 2: 1+						
		• TNFR						
		• Interleukin 6						
		Model 3: 2+						
		• Serial						
		measurement of						
		cytokines						

Study	Derivation	Model/ Variables	Selection	Linear	Over-	Model	Calibration	Discrimination
	Validation			Gradient	fitting	assumptions		(c-statistic)
O'Connor	Derivation	HF-ACTION:	Based on	Checked	No	n.r.	Correlation	Overall=0.73
2012 [12]		Exercise duration	univariable	by			(r=0.99 at	
		• Urea	analysis	restrictive			1,2 and 3	
		• Sex		cubic			years and	
		• BMI		spline			0.98 at 5	
							years)	
Herrmann	Derivation	• Peak VO ₂	Based on	n.r.	Yes	n.r.	n.r.	† Overall=0.91
2012 [13]		<14ml/kg/min	previous		(31			
		• Uric acid	knowledge		deaths			
		>565µmol/L			and 5			
		• LVEF<22%			variables			
		• Cholesterol			tested)			
		<5.27mmol/L						
		• sTNF-R1						
		>1016pg/L						

Study	Derivation	Model/ Variables	Selection	Linear	Over-	Model	Calibration	Discrimination
	Validation			Gradient	fitting	assumptions		(c-statistic)
Scrutinio	Derivation	• Age	Based on	n.r.	No	n.r.	H-L test	Overall=0.74
2012 [14]		• Ischemic CMP	univariable				(p>0.45)	
		• Anemia	analysis					
		• LVEF						
		• Renal function						
Pocock	Derivation	• Age	Based on	n.r.	No	n.r.	Graphically	n.r.
2012 [15]		• Gender	statistical				observed vs.	
		• BMI	significance				predicted =	
		• Current smoker					Adequate	
		• Systolic BP						
		• Diabetes						
		• NYHA class						
		• LVEF						
		• COPD						
		• HF duration						

Creatinine			
• β-blockers			
• ACE-I/ARB			

‡ Authors conducted a subgroup analysis based on underlying etiology and LVEF and reported that c-index was equal in ischemic, non-ischemic CMP and patients with LVEF <30%, but lower (c-statistic = 0.73) in patients with LVEF ≥30%.

† Authors reported that a model excluding cholesterol has similar c-statistic and that a model including uric acid, sTNF-R1, LVEF and NYHA class (<3) instead of peak VO₂ had an overall c-statistic of 0.84.

LVEF, left ventricular ejection fraction; VO2, oxygen consumption; CT, cardio-thoracic; VT, ventricular taqui-arrhythmia; LVH, left ventricular hypertrophy; ECG, electro-cardiogram; SDNN, standard deviation of all R-to-R intervals on 24-h; MRT, mean response time; BP, blood pressure; CVA, cerebro-vascular accident; NYHA, New York Heart Association; BMI, body mass index; BBB, bundle branch block; MI, myocardial infarction; PVD, peripheral vascular disease; ICD, internal cardiac defibrillator; MFH; metabolic, functional, hemodynamic; CPX, cardiopulmonary exercise test; MRT, mean response time; MI; myocardial infarction; DSC, Dyssynchrony, posterolateral Scar location and Creatinine; CRT, cardiac resynchronization therapy; CV, cardiovascular; BNP, brain natriuretic peptide; COPD, chronic obstructive pulmonary disease; CMP, cardiomyopathy; sTNF-R1, soluble tumor necrosis factor alpha receptor 1; H-L, Hosmer and Lemeshow; ACE-I, angiotensin converting enzyme inhibitor; ARB, angiotensin receptor blocker; n.r., not reported; n/a, not applicable.

^{*} This model was validated by bootstrapping but discrimination capacity on bootstrapping is not reported.

References of Supplemental tables 4, 5 and 6

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