

REVIEW



Mortality prediction models in the adult critically ill: A scoping review

Britt E. Keuning¹  | Thomas Kaufmann²  | Renske Wiersema¹  |
Anders Granholm³  | Ville Pettilä⁴  | Morten Hylander Møller^{3,5}  |
Christian Fynbo Christiansen⁶  | José Castela Forte^{1,7}  | Harold Snieder⁸  |
Frederik Keus¹  | Rick G. Pleijhuis⁹  | Iwan C. C. van der Horst^{1,10}  |

HEALICS consortium

¹Department of Critical Care, University of Groningen, University Medical Center Groningen, Groningen, The Netherlands

²Department of Anesthesiology, University of Groningen, University Medical Center Groningen, Groningen, The Netherlands

³Department of Intensive Care, Copenhagen University Hospital, Rigshospitalet, Copenhagen, Denmark

⁴Division of Intensive Care Medicine, Department of Anesthesiology, Intensive Care and Pain Medicine, University of Helsinki and Helsinki University Hospital, Helsinki, Finland

⁵Centre for Research in Intensive Care, Copenhagen University Hospital, Rigshospitalet, Copenhagen, Denmark

⁶Department of Clinical Epidemiology, Aarhus University Hospital, Aarhus, Denmark

⁷Bernoulli Institute for Mathematics, Computer Science and Artificial Intelligence, University of Groningen, Groningen, The Netherlands

⁸Department of Epidemiology, University of Groningen, University Medical Center Groningen, Groningen, The Netherlands

⁹Department of Internal Medicine, University of Groningen, University Medical Center Groningen, Groningen, The Netherlands

¹⁰Department of Intensive Care, Maastricht University Medical Center+, Maastricht University, Maastricht, The Netherlands

Correspondence

Iwan C. C. van der Horst, Department of Intensive Care, Maastricht University Medical Center+, Maastricht University, Maastricht, The Netherlands.
Email: iwan.vander.horst@mumc.nl

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Background: Mortality prediction models are applied in the intensive care unit (ICU) to stratify patients into different risk categories and to facilitate benchmarking. To ensure that the correct prediction models are applied for these purposes, the best performing models must be identified. As a first step, we aimed to establish a systematic review of mortality prediction models in critically ill patients.

Methods: Mortality prediction models were searched in four databases using the following criteria: developed for use in adult ICU patients in high-income countries, with mortality as primary or secondary outcome. Characteristics and performance measures of the models were summarized. Performance was presented in terms of discrimination, calibration and overall performance measures presented in the original publication.

Results: In total, 43 mortality prediction models were included in the final analysis. In all, 15 models were only internally validated (35%), 13 externally (30%) and 10 (23%) were both internally and externally validated by the original researchers. Discrimination was assessed in 42 models (98%). Commonly used calibration measures were the Hosmer-Lemeshow test (60%) and the calibration plot (28%).

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Calibration was not assessed in 11 models (26%). Overall performance was assessed in the Brier score (19%) and the Nagelkerke's R^2 (4.7%).

Conclusions: Mortality prediction models have varying methodology, and validation and performance of individual models differ. External validation by the original researchers is often lacking and head-to-head comparisons are urgently needed to identify the best performing mortality prediction models for guiding clinical care and research in different settings and populations.

1 | INTRODUCTION

Outcome prediction models, severity scales and risk scores are prognostic tools to estimate the probability for a pre-specified outcome.¹ These prognostic tools use variables (eg about the severity of illness) to predict outcome, often mortality, in a specific patient population such as the critically ill. In the intensive care unit (ICU), mortality prediction models may be applied to stratify patients in different risk categories and to facilitate benchmarking using standardized mortality rates. An accurate mortality prediction model provides a stratification of the risk of an outcome at a population level. These models generally provide a numerical estimate of that risk based on estimates from previous populations.² Per definition, all mortality prediction models are best suited for use at a population level and not for individual prognostication, as uncertainty for individual patients remains high.^{3,4}

Several models are widely known and broadly applied such as the Acute Physiology and Chronic Health Evaluation (APACHE) I-IV, the Mortality Prediction Model (MPM) and the Simplified Acute Physiology Score (SAPS) I-III,⁵ whereas others like the Intensive Care National Audit & Research Centre (ICNARC) are used solely in one country.⁶ Previous literature has only reviewed commonly used models, models with different outcome than mortality or disease- or organ-specific prognostic models.^{3-5,7,8} To the best of our knowledge, no study has systematically assessed which mortality prediction models have been developed and validated for broad cohorts of adult critically ill patients.

1.1 | Rationale and objective

The objective of this study was to provide an overview of available mortality prediction models in adult critically ill patients as a step-up towards future head-to-head comparison of model performance through systematic external validation.

2 | METHODS

2.1 | Protocol and registration

This scoping review was performed following our protocol (Appendix S1) and was reported in accordance with the PRISMA-ScR checklist.⁹

Editorial Comment

In this review, mortality prediction models in intensive care have been identified. Characteristics and performance of 43 individual models are summarized according to documentation in the original publications so that validation and predictive performances can be compared.

Notably, we aimed to publish the protocol on PROSPERO, but during the process it showed that PROSPERO currently does not accept registrations for scoping reviews, literature reviews or mapping reviews.

2.2 | Search strategy

We conducted a systematic search of MEDLINE, EMBASE, Web of Science and The Cochrane Central Register of Controlled Trials (CENTRAL) to identify relevant ICU mortality prediction models (Appendix S1). Mortality was chosen as the outcome of interest, as prediction models were originally developed to identify patients with high mortality risk. For all databases, except the CENTRAL database, the search period encompassed a period starting from the 1st January 2008 to the 21st April 2019. We used snowballing, that is, searching references and related articles, to identify additional prediction models that were published before 2008.

One author ran the search, after which the screening of records and data extraction were performed in duplicate. All records were screened based on title and/or abstract. Papers clearly irrelevant to the purpose were excluded. The remaining articles were screened for eligibility. Consulting a third opinion solved disagreements. More detailed information is presented in the protocol (Appendix S1).

2.3 | Eligibility criteria

To be considered eligible, mortality prediction models had to meet the following criteria: (a) originally developed specifically for use in adult critically ill patients as defined by the included studies, (b) representing

broad groups of ICU patients (with large diversity of admission diagnoses, eg non-diabetic patients, medical admissions, surgical admissions, etc), (c) availability of the original article in English and (d) mortality at any time as (primary or secondary) outcome of interest.

Prediction models were excluded (a) when developed for low- or middle-income countries, as characteristics of ICU patients in these countries often substantially differ from those in high-income countries and, epidemiological data from low-income countries have been frequently unavailable,^{10,11} (b) when developed as a digital model or derived from a machine-learning algorithm, since code and data availability are not requirements in all journals. Since our utmost goal is to make a head-to-head comparison of available mortality prediction models using an independent external validation cohort, the code or data necessary to retrieve the underlying prediction model formula are required to reproduce the prediction models. (c) When the development of multiple customized prediction models was described in one article, but no final model was proposed, the prediction models were excluded. Finally, (d) we excluded prediction models specifically developed for subgroups of intensive care patients such as those with sepsis, trauma, cardiac and neurological patients. Studies not specifying inclusion of these subgroups within a wider, general ICU population were considered to be eligible. Prediction models developed in a medical or surgical ICU were included.

2.4 | Data extraction

If multiple mortality outcomes (eg at different time points) were used, we used the primary outcome in the original publication (or the first mortality outcome if the primary outcome was not mortality) to describe the performance of the prediction model.

Details on the development process of the mortality prediction models included were shown, as well as the number of variables included in the prediction models, mortality rate in each development setting and method of handling of missing data. To give an overview of the performance of all mortality prediction models, for example, values from discrimination, calibration and overall performances measures¹² for mortality were presented for development and internal or external validation cohorts in the original publication (if available).

The discrimination measure presented was the C-statistic (area under the receiver operating characteristic curve [AUROC]), calibration measures presented were goodness-of-fit tests like the Hosmer-Lemeshow (HL) test, calibration plot and calibration slope, and the overall performance measures presented were the Nagelkerke's R^2 and the Brier score.¹²

Preferable values from external validation were presented if both internal and external validation values were present in the

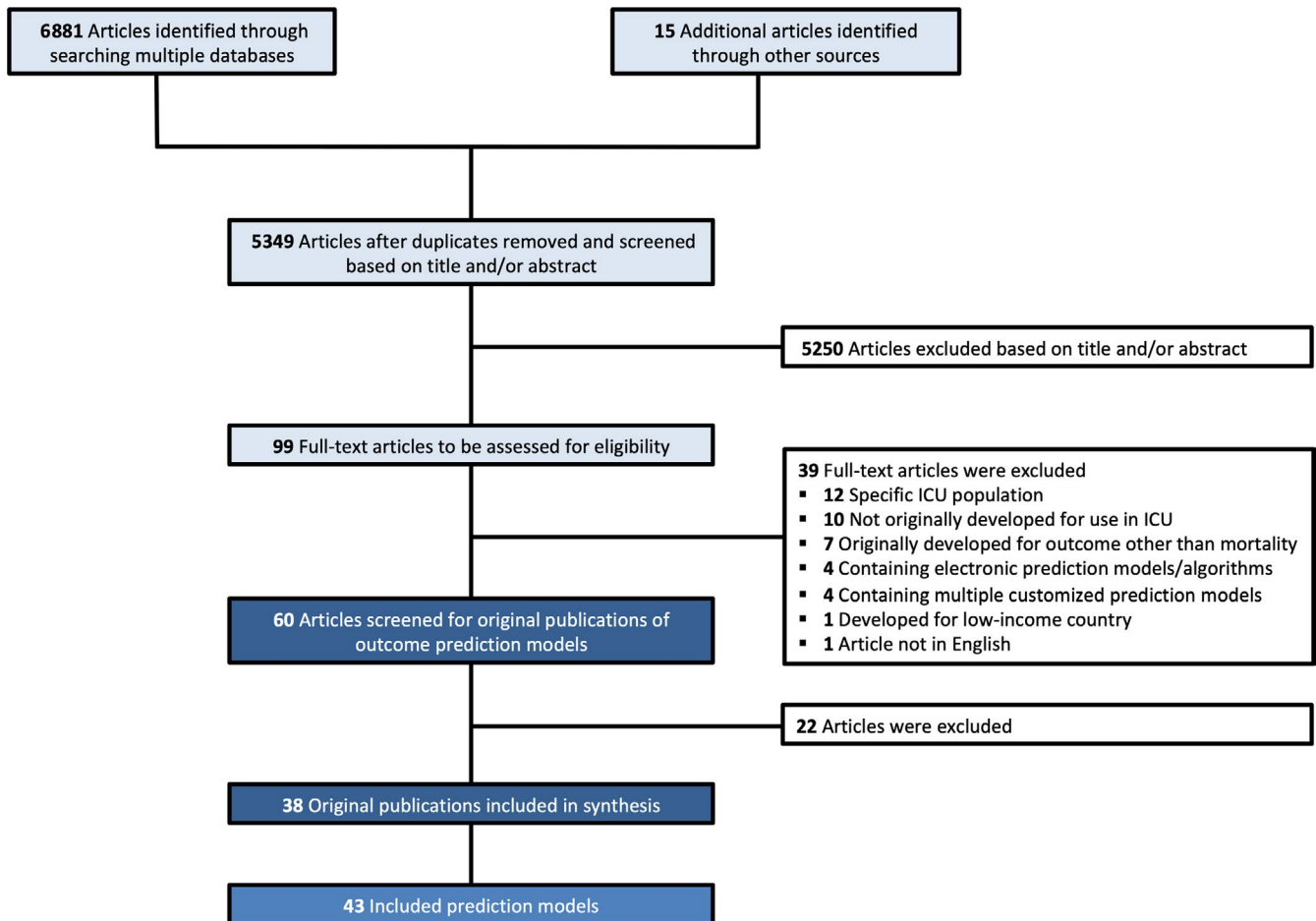


FIGURE 1 Flow diagram of the search [Colour figure can be viewed at wileyonlinelibrary.com]

original publication. If not available, values of internal validation cohorts were presented. External validation was defined as using a separate individual dataset for validation of the mortality prediction model (ie no split sampling of a dataset also used for the development of the model).

Citations of original publications were screened for internal and/or external validation articles and shown as being present (+) or absent (-). A list of variables sought for in the identified articles can be found in Appendix S1.

3 | RESULTS

The selection of sources of evidence can be found in the flowchart (Figure 1). Articles evidently developed for specific groups of patients (ie sepsis, trauma, cardiac, neurological patients) were excluded based on the title and/or abstract. Evaluating 99 full-text articles for eligibility resulted in exclusion of another 39 articles, leaving 60 articles that were screened for original publications. Eventually, 43 relevant mortality prediction models reported in 38 publications were extracted and included in the final analysis.

3.1 | Characteristics of the included mortality prediction models

Characteristics of the mortality prediction models and underlying derivation cohorts are presented in Table 1. In all, 19 mortality prediction models (44%) were developed using prospectively collected data specifically gathered for the development of the prediction model,^{6,13-27} whereas 24 (56%) were developed using either retrospective data²⁸⁻⁴⁴ or prospective data previously collected for other purposes.⁴⁵⁻⁴⁹ The start of data collection for the development cohorts spanned 36 years (1979-2015), and the duration of the cohort studies varying from 2 months up to 10 years for each cohort. Two mortality prediction models (4.7%) did not report the timespan during which their development cohort was assembled.^{22,33} In all, 31 mortality prediction models (74%) were developed in a single country,^{14,18-27,29,31,33-45,47,49} six (14%) in neighbouring countries (two or more)^{6,13,28,30,32,46} and five (12%) were developed in multiple countries worldwide.^{15-17,48} The number of patients included in the development databases ranged from 232 to 731 611 patients with a median of 4,895 (IQR 528-35 878). The minimum age at which patients were included was 15 years (2.3%).³⁵ In all, 11 mortality prediction models (26%) did not specify age.^{6,13,23,25,29,31,36,38,42,46} The number of variables included in the mortality prediction models varied from 5 up to 5695, with a median of 16 (IQR 9-24).

3.2 | Outcome measures

The timing of mortality outcome varied between the studies. Hospital mortality was the most frequently used

primary outcome in 29 (67%) mortality prediction models.^{6,13-19,21,22,24,27,28,30-33,35,36,38,41-43,45,46} Other primary outcome variables were ICU mortality (7%),^{23,26,34} 28-day mortality (4.7%),^{39,44} 90-day mortality (4.7%),^{48,49} 3- to 28-day mortality (4.7%),⁴⁰ 30-day mortality (2.3%),⁴⁷ 180-day mortality (2.3%),²⁰ 6-month mortality (2.3%),²⁵ 15-year mortality (2.3%),³⁷ and 6- and 12-month mortality (2.3%).²⁹

Secondary outcomes were 1-month mortality after ICU admission (4.7%),^{24,31} hospital mortality (4.7%),^{29,34} ICU mortality (2.3%),⁴⁵ 3-month mortality after ICU admission (2.3%),³¹ 6-month mortality after ICU admission (2.3%),³¹ 9-month mortality (2.3%),⁴⁷ 1-year mortality (2.3%)⁴⁵ and length of stay (2.3%).²⁴ Of the 43, 37 mortality prediction models (86%) did not prognosticate any secondary outcome.^{6,13-23,25-28,30,32,33,35-44,46,48,49}

Hospital mortality rates of the development cohorts varied from 6.9% to 48% and were not reported for nine mortality prediction models (21%).^{6,15,18,29,33,40,42}

For 21 mortality prediction models (49% of 43), data were collected within the first 24 hours after patient admission to the ICU.^{6,13,14,17-19,24,26,27,30,31,34,38,39,42,44,47-49} For 11 prediction models (26%), data on ICU admission were collected,^{16,23,25,28,32,35,36,41,43,45,46} whereas for the remaining prediction models data timing varied from 24 days before admission up to 5 days after patient admission to the ICU.

Handling of missing data was not reported in 11 mortality prediction models (26%),^{23,25,26,31,33,38,39,41,45,46,49} 20 prediction models (47% of 43) excluded records with missing data,^{6,14,16,19,21,24,27,28,30,32,34,40,42-44} six prediction models (14%) imputed values with normal or mean values^{15,17,18,20,22,29} and four prediction models (9.3%) reported no missing data.^{13,35-37} The remaining two prediction models (4.7%) excluded patients when more than a certain percentage of the data was missing (>5% or >25%).^{47,48}

3.3 | Discrimination, calibration and overall performance measures

Discrimination, calibration and overall performance measures are presented in Table 2. Of the 43 mortality prediction models, 15 (35%) were only internally validated,^{23,26,28-31,33,38-41,44,46,48} 13 (30%) only externally,^{16,19-21,25,35,36,42,43,47} 10 (23%) were both internally and externally validated,^{6,13-15,17,18,22,32,34,37} and 5 prediction models (12%) were not validated at all.^{24,27,45,49} In all, 15 prediction models (35%) included a description of an external validation in their original publication.^{13,16,20-22,25,34-36,42,43,47}

Discrimination was expressed as the AUROC in 42 of the 43 mortality prediction models original publications (98%). Only the APACHE II model did not report an AUROC value in the original publication.¹⁹ In the development cohorts, the lowest discrimination was AUROC 0.72 (95% CI 0.71-0.74),⁴⁸ and the highest AUROC 0.91 (95% CI not specified).³⁰ In the validation cohorts, the lowest AUROC was 0.58 (95% CI not specified),⁴⁴ and the highest AUROC 0.95 (0.91-0.99).²³

Calibration measures were expressed by various statistical measures. The HL goodness-of-fit test was used in 26 mortality prediction

TABLE 1 Characteristics of the development of the 43 mortality prediction models

Mortality prediction model	Year published	Development database	Cohort assembly period	ICU population	Number of variables ^a	Outcome		Handling of missing data	
						Primary	Secondary		
ICNARC Harrison et al ⁶	2007	216 626 Prospective	December 1995–August 2003	General, adult patients in England, Wales and Ireland	16	Hospital mortality	–	Worst values and total urine output in initial 24 h in ICU	Exclusion
ICNARC-II Ferrando-Vivas et al ¹³	2017	155 239 Prospective	01/01/2012– 31/12/2012	General, adult patients in England, Wales and Ireland	23	Hospital mortality	–	Worst values and total urine output in initial 24 h in ICU	No missing data
APACHE IV Zimmerman et al ⁴	2006	66 270 Prospective	01/01/2002– 31/12/2003	General, adult (≥16 y) patients in the USA	142	Hospital mortality	–	Worst values in initial 24 h in ICU	Exclusion
SAPS III Moreno et al ¹⁵	2005	13 428 ^b Prospective	14/10/2002– 15/12/2002	General, adult (≥16 y) patients worldwide	20	Hospital mortality	–	ICU admission ± 1 h	Imputation of normal values
MPM ₀ -II Lemeshow et al ¹⁶	1993	12 610 Prospective	17/04/1989– 31/07/1990 (dataset I) and 30/09/1991– 27/12/1991 (dataset II)	General, adult (≥18 y) patients in Europe and the USA	15	Hospital mortality	–	ICU admission	Exclusion
MPM ₂₄ -II Lemeshow et al ^{16,21}	1993	10 357 Prospective	17/04/1989– 31/07/1990 (dataset I) and 30/09/1991– 27/12/1991 (dataset II)	General, adult (≥18 y) patients in Europe and the USA	13	Hospital mortality	–	At 24 h in ICU	Exclusion
SAPS II Le Gall et al ¹⁷	1993	8369 Prospective	30/09/1991– 28/02/1992	General, adult (≥18 y) patients in Europe and North-America	17	Hospital mortality	–	Worst values in initial 24 h in ICU	Imputation of normal values
APACHE III Knaus et al ¹⁸	1991	7848 ^b Prospective	May 1988–November 1989	General, adult (≥16 y) patients in the USA	26	Hospital mortality	–	Worst values in initial 24 h in ICU	Imputation of normal values
APACHE II Knaus et al ¹⁹	1985	5030 Prospective	1979–1982	General, adult (≥16 y) patients in the USA	18	Hospital mortality	–	Worst values in initial 24 h in ICU	Exclusion

(Continues)

TABLE 1 (Continued)

Mortality prediction model	Year published	Development database	Cohort assembly period	ICU population	Number of variables ^a	Outcome		Data collection	Handling of missing data
						Primary	Secondary		
SUPPORT Knaus et al ²⁰	1995	4301 Prospective	June 1989-June 1991	General, adult (≥18 y) patients in the USA	15	180-day mortality	-	2072/4301 (48.2%)	Imputation of normal values, missing data at day 3 were imputed with day 1 values
MPM ₄₈ -II Lemeshow et al ²¹	1994	2049 Prospective	17/04/1989- 31/07/1990	General, adult (≥18 y) patients in the USA	13	Hospital mortality	-	307/2049 ^b (15.0%)	Exclusion
MPM ₇₂ -II Lemeshow et al ²¹	1994	1497 Prospective	17/04/1989- 31/07/1990	General, adult (≥18 y) patients in the USA	13	Hospital mortality	-	418/1497 ^b (27.9%)	Exclusion
TRIOS Timsit et al ²²	2001	893 Prospective	Not reported (valida- tion dataset in March 1999)	General, adult (≥16 y) patients, hospitalized >48 h in France	32	Hospital mortality	-	268/893 (30.0%)	Imputation of normal values
Mortality Risk Score Dólera-Moreno et al ²³	2016	844 Prospective	January 2013-April 2014	General, adult patients in Spain	6	ICU mortality	-	91/844 (10.8%)	Not reported
Mortality Multifactor Model Li et al ²⁴	2017	500 Prospective	01/03/2014- 30/04/2014	General, adult (≥18 y) patients in China	36	Hospital mortality	Mortality 30 days after ICU admis- sion, LOS	102/500 (20.4%)	Exclusion
Mortality Prognostic Model Hadique et al ²⁵	2017	500 Prospective	November 2013-April 2014	Medical, adult pa- tients in the USA	44	6-month mortality	-	180/500 (36.0%)	Not reported
Mortality Prediction Model Fika et al ²⁶	2018	400 Prospective	January 2012-July 2013	General, adult (≥18 y) patients in Greece	12	ICU mortality	-	131/400 (23.8%)	Not reported
APACHE II-APM Nematifard et al ²⁷	2018	304 Prospective	June 2014-November 2016	General, adult (≥16 y) patients in Iran	19	Hospital mortality	-	96/304 (31.6%)	Exclusion

(Continues)

TABLE 1 (Continued)

Mortality prediction model	Year published	Development database	Cohort assembly period	ICU population	Number of variables ^a	Outcome		Hospital mortality rate in each development setting	Data collection	Handling of missing data
						Primary	Secondary			
APACHE III-APM Nematifard et al ²⁷	2018	304 Prospective	June 2014-November 2016	General, adult (≥16 y) patients in Iran	27	Hospital mortality	-	96/304 (31.6%)	Worst values in initial 24 h in ICU	Exclusion
ANZROD0 Paul et al ²⁸	2017	731 611 Retrospective	01/01/2006-31/12/2015	General, adult (≥16 y) patients in Australia and New Zealand	11	Hospital mortality	-	69 503/731 611 ^b (9.5%)	ICU admission	Exclusion
MMI Min et al ²⁹	2017	354 154 ^b Retrospective	January 2003-December 2013	Medical, veteran ICU patients in the USA	5695	All-cause mortality at 6- and 12-months post-hospital discharge	Hospital mortality	Not reported	Worst values of 24 h before and 24 h after admission	Imputation of mean values
ANZROD Paul et al ³⁰	2013	304 149 Retrospective	01/01/2004-31/12/2009	General, adult (≥16 y) patients in Australia and New Zealand	38	Hospital mortality	-	34 369 ^b (11.3%)	Worst values in initial 24 h in ICU	Exclusion
Customized APACHE IV Brinkman et al ³¹	2013	77 616 Retrospective	01/01/2008-01/07/2011	Non-CABG, adult critically ill patients in the Netherlands	142	Hospital mortality	Mortality at 1, 3 and 6 months after ICU admission	12 186/77 616 ^b (15.7%)	First 24 h in ICU	Not reported
MPM ₀ -III Higgins et al ³²	2005	74 578 Retrospective	October 2001-March 2004	General, adult (≥18 y) patients in the USA, Canada and Brazil	16	Hospital mortality	-	10 292/74 578 (13.8%)	ICU admission	Exclusion
NQF-ICOMmort Philip R. Lee Institute ³³	2016	40 395 Retrospective	Not reported	General, adult (≥18 y) patients in the USA	17	Hospital mortality	-	Not reported	1 h prior to ICU admission to 1 h after admission	Not reported
OASIS Johnson et al ³⁴	2013	39 070 Retrospective	01/01/2007-15/09/2011	General, adult (≥16 y) patients in the USA	10	ICU mortality	Hospital mortality	4571/39 070 ^b (11.7%)	Worst values and total urine output in initial 24 h in ICU	Exclusion
COPE-4 Duke et al ³⁵	2013	35 878 Retrospective	01/07/2004-30/06/2006	General, adult (≥15 y) patients in Australia	6	Hospital mortality	-	4415/35 878 (12.3%)	ICU admission (mechanical ventilation during ICU admission)	No missing data

(Continues)

TABLE 1 (Continued)

Mortality prediction model	Year published	Development database	Cohort assembly period	ICU population	Number of variables ^a	Outcome			Handling of missing data	
						Primary	Secondary	Hospital mortality rate in each development setting		Data collection
RDW-SAPS Hunziker et al ⁴⁵	2012	17 922 Retrospective	January 2001-December 2008	General, adult (≥18 y) patients in the USA	15	Hospital mortality	ICU mortality, 1-year mortality	2007/17 922 ^b (11.2%)	ICU admission	Not reported
COPE Duke et al ³⁶	2008	17 880 Retrospective	01/07/2004-30/06/2005	General, adult patients in Australia	5	Hospital mortality	-	2186/17 880 (12.1%)	ICU admission (mechanical ventilation during ICU admission)	No missing data
PREDICT Ho et al ³⁷	2008	11 930 Retrospective	1989-2002	General, adult (≥16 y) patients in Australia	6	15-year mortality	-	829/11 930 ^b (6.9%)	First 5 days in ICU	No missing data
High-Risk Selection System Iapichino et al ⁴⁶	2006	8248 Retrospective	October 1994-February 1995	General, adult patients (>24 h in ICU) in Europe	16	Hospital mortality	-	1617/8248 ^b (19.6%)	ICU admission	Not reported
GV-SAPS II Liu et al ⁴⁷	2016	4895 Retrospective	2001-2008	Non-diabetic, adult (≥18 y) patients in the USA	20	30-day mortality	9-month mortality	649/4895 (13.3%)	First 24 h in ICU	When >5% exclusion, <5% not reported
MODS/NEMS Kao et al ³⁸	2016	4321 Retrospective	01/01/2009-30/11/2012	General, adult patients in Canada	32	Hospital mortality	-	986/4321 (22.8%)	First 24 h in ICU	Not reported
SMS-ICU Granholm et al ⁴⁸	2018	4086 Retrospective	23/12/2009-30/06/2016	General, adult (≥18 y), acutely admitted patients worldwide	7	90-day mortality	-	1403/4086 (34.3%)	Worst values in initial 24 h in ICU	Multiple imputations, exclusion when >25%
P- model Umegaki et al ³⁹	2010	3505 Retrospective	01/01/2007-31/12/2007	General, adult (≥20 y) patients in Japan	10	Mortality at 28 days after the first ICU day	-	336/3505 ^b (9.6%)	First 24 h in ICU	Not reported
BCV model Huang et al ⁴⁰	2013	1624 Retrospective	01/01/2006-01/12/2008	General, adult (≥18 y) patients in Taiwan	6	Daily probability of mortality from day 3 to day 28 post-ICU admission	-	Not reported	Daily complete blood count	Exclusion

(Continues)

TABLE 1 (Continued)

Mortality prediction model	Year published	Development database	Cohort assembly period	ICU population	Number of variables ^a	Outcome		Hospital mortality rate in each development setting	Data collection	Handling of missing data
						Primary	Secondary			
BCV/APACHE II model Huang et al ⁴⁰	2013	1624 Retrospective	01/01/2006-01/12/2008	General, adult (≥18 y) patients in Taiwan	24	Daily probability of mortality from day 3 to day 28 post-ICU admission	-	Not reported	Daily complete blood count, APACHE II score in the first 24 h in ICU	Exclusion
CREEK Stachon et al ⁴¹	2008	528 Retrospective	April 2003-January 2004	Medical, adult (≥18 y) patients in Germany	8	Hospital mortality	-	87/528 (16.5%)	ICU admission	Not reported
SAPS-R Viviani et al ⁴²	1991	351 Retrospective	01/01/1986-31/10/1988	General, adult patients in France	5	Hospital mortality	-	Not reported	Worst values in initial 24 h in ICU	Exclusion
SAPS-E Viviani et al ⁴²	1991	351 Retrospective	01/01/1986-31/10/1988	General, adult patients in France	7	Hospital mortality	-	Not reported	Worst values in initial 24 h in ICU	Exclusion
25OHD Deyo-Charlson Comorbidity Index Mahato et al ⁴⁹	2016	310 Retrospective	01/06/2012-30/05/2015	General, adult (≥18 y) patients in the USA	18	90-day mortality after ICU admission	-	59/310 (19.0%)	First 24 h in ICU	Not reported
DELAWARE Stachon et al ⁴³	2008	271 Retrospective	April 2003-January 2004	Surgical, adult (≥18 y) patients in Germany	9	Hospital mortality	-	67/271 (24.7%)	ICU admission	Exclusion
Simplified Mortality Score Goag et al ⁴⁴	2018	232 Retrospective	June 2015-February 2016	Medical, adult (≥18 y) patients in Korea	8	28-day mortality	-	72/232 ^b (31.1%)	Within 24 h of ICU admission	Exclusion

Abbreviations: ANZROD, Australian and New Zealand Risk Of Death; APACHE, Acute Physiology and Chronic Health Evaluation; APM, adductor pollicis muscle; BCV, blood cell variability; COPE, critical care outcome prediction equation; CREEK, critical risk evaluation by early keys; DELAWARE, Dense Laboratory Whole Blood Applied Risk Estimation; GV, glucose variability; ICNARC, Intensive Care National Audit Research Centre; ICU, intensive care unit; LOS, length of stay; MMI, multi-morbidity index; MODS, multiple organs dysfunction score; MPM; mortality prediction model; NEMS, nine equivalents nursing manpower use score; NQF-ICOMmort, national quality forum ICU outcomes model (mortality); OASIS, Oxford acute severity of illness score; PREDICT, predicted risk, existing diseases and intensive care therapy; RDW, red cell distribution width; SAPS, simplified acute physiology score; SMS-ICU, simplified mortality score for the intensive care unit; SQ, surprise question; SUPPORT, study to understand prognoses and preferences for outcomes and risks of treatments; TRIOS, three-day recalibrating ICU outcomes.

^aWhen (parts of) other mortality prediction models were used as variables in a mortality prediction model (eg the Charlson Comorbidity Index and APACHE III as variable in the Mortality Prognostic Model), variables included in these specific mortality prediction models were also taken into account.

^bEstimated based on information in original publication.

TABLE 2 Performance of the 43 mortality prediction models

Mortality prediction model	Validated? ^a		AUROC (95% CI) Development cohort ^b	Calibration Development cohort ^b	Overall performance Development cohort ^b	Type of validation cohort in original publication	AUROC (95% CI) Validation cohort	Calibration Validation cohort	Overall performance Validation cohort
	Internally	Externally							
ICNARC Harrison et al ⁶	+	+	-	-	-	Internal validation dataset	0.87 (n.s.)	-	Brier score: 0.132
ICNARC-II Ferrando-Vivas et al ¹³	+	+	0.89 (0.89-0.89)	-	Brier score: 0.103	External validation dataset	0.89 (0.88-0.89)	Calibration plot present	Brier score: 0.108
APACHE IV Zimmerman et al ¹⁴	+	+	-	-	-	External validation dataset	0.88 (n.s.)	HL X ² : 16.8 (P = .08)	-
SAPS III Moreno et al ¹⁵	+	+	-	-	-	Internal validation dataset	0.85 (n.s.)	HL H-statistic: 10.6 (P = .39) HL C-statistic: 14.3 (P = .16) Calibration plot present	-
MPM ₀ -II Lemeshow et al ¹⁶	-	+	0.84 (n.s.)	HL C-statistic: 6.2 (P = .62)	-	External validation dataset	0.82 (n.s.)	HL C-statistic: n.s. (P = .33)	-
MPM ₂₄ -II Lemeshow et al ^{16,21}	-	+	0.84 (n.s.)	HL C-statistic: 4.9 (P = .76)	-	External validation dataset	0.84 (n.s.)	HL C-statistic: 12.9 (P = .23)	-
SAPS II Le Gall et al ¹⁷	+	+	0.88 (0.87-0.90)	HL H-statistic: 3.70 (P = .88)	-	Internal validation dataset	0.86 (0.84-0.88)	HL H-statistic: n.s. (P = .10)	-
APACHE III Knaus et al ¹⁸	+	+	-	-	-	Internal validation dataset	0.90 (n.s.) ^c	-	-
APACHE II Knaus et al ¹⁹	-	+	-	-	-	-	-	-	-
SUPPORT Knaus et al ²⁰	-	+	0.79 (n.s.)	-	-	External validation dataset	0.78 (n.s.)	Calibration plot present	-
MPM ₄₈ -II Lemeshow et al ²¹	-	+	0.81 (n.s.)	HL C-statistic: 11.7 (P = .31)	-	External validation dataset	0.80 (n.s.)	HL C-statistic: 8.4 (P = .59)	-
MPM ₁₂ -II Lemeshow et al ²¹	-	+	0.79 (n.s.)	HL C-statistic: 11.6 (P = .31)	-	External validation dataset	0.75 (n.s.)	HL C-statistic: 10.4 (P = .41)	-

(Continues)

TABLE 2 (Continued)

Mortality prediction model	Validated? ^a		AUROC (95% CI) Development cohort ^b	Calibration Development cohort ^b	Overall performance Development cohort ^b	Type of validation cohort in original publication	AUROC (95% CI) Validation cohort	Calibration Validation cohort	Overall performance Validation cohort
	Internally	Externally							
TRIOS Timsit et al ²²	+	+	0.79 (0.77-0.82)	HL C-statistic: 5.6 (P = .70)	-	External validation dataset	0.83 (0.78-0.87)	-	-
Mortality Risk Score Dólera-Moreno et al ²³	+	-	-	-	-	Internal validation dataset	0.95 (0.91-0.99)	Likelihood ratio test X ² : 296.8 ^c	-
Mortality Multifactor Model Li et al ²⁴	-	-	0.84 (0.80-0.87)	HL X ² : 12.3 (P = .14) Calibration plot present	-	-	-	-	-
Mortality Prognostic Model Hadique et al ²⁵	-	+	0.83 (0.80-0.87)	HL statistic: 6.5 (P = .59)	-	External validation dataset	0.84 (0.81-0.88)	HL statistic: 9.2 (P = .33)	-
Mortality Prediction Model Fika et al ²⁶	+	-	-	-	-	Internal validation dataset	0.85 (0.73-0.97)	HL X ² : 4.9 (P = .77)	-
APACHE II-APM Nematifard et al ²⁷	-	-	0.85 (0.81-0.90)	-	-	-	-	-	-
APACHE III-APM Nematifard et al ²⁷	-	-	0.87 (0.82-0.91)	-	-	-	-	-	-
ANZROD0 Paul et al ²⁸	+	-	0.85 (0.85-0.86)	HL C-statistic: 459.3	Brier score: 0.069 Adjusted Brier score: 0.196	Internal validation dataset	0.85 (0.85-0.85)	HL C-statistic: 264.9 Calibration plot present	Brier score: 0.069 Adjusted Brier score: 0.190
MIMI Min et al ²⁹	+	-	-	-	-	Internal validation dataset	6-month mortality: 0.86 (0.85-0.86) 12-month mortality: 0.84 (0.83-0.84)	-	Brier score: 0.21 ^{c,d}

(Continues)

TABLE 2 (Continued)

Mortality prediction model	Validated? ^a		AUROC (95% CI) Development cohort ^b	Calibration Development cohort ^b	Overall performance Development cohort ^b	Type of validation cohort in original publication	AUROC (95% CI) Validation cohort	Calibration Validation cohort	Overall performance Validation cohort
	Internally	Externally							
ANZROD Paul et al ³⁰	+	-	0.91 (n.s.)	HL C-statistic: 189.5 HL H-statistic: 174.1 Cox calibration regression slope: 1	Brier score: 0.065	Internal validation dataset	0.90 (n.s.)	HL C-statistic: 104.9 HL H-statistic: 111.4 Cox calibration regression slope: 0.98 Calibration plot present	Brier score: 0.066
Customized APACHE IV Brinkman et al ³¹	+	-	0.88 (0.88-0.88)	Calibration plot present	Brier score: 0.09	Internal validation dataset	-	-	-
MPM ₀ -III Higgins et al ³²	+	+	0.83 (0.82-0.83)	HL statistic: 11.5 (P = .17)	-	Internal validation dataset	0.82 (0.82-0.83)	HL statistic: 11.6 (P = .31)	-
NQF-ICOMmort Philip R. Lee Institute ³³	+	-	-	-	-	Internal validation dataset	0.82 (0.81-0.83)	HL C statistic: 12.0 (P = .28) HL H statistic: 16.9 (P = .08) Calibration plot present	-
OASIS Johnson et al ³⁴	+	+	-	-	-	External validation dataset	0.90 (P < .0003) ^e	HL X ² : 19.6 ^e	Brier score: 0.048 ^e
COPE-4 Duke et al ³⁵	-	+	-	-	-	External validation dataset	-(0.82-0.83)	HL H-statistic: 14.8 (P = .06) Correlation of calibration plot R ² : 0.99 Calibration plot present	-
RDW-SAPS Hunziker et al ⁵⁰	-	-	0.77 (n.s.)	Quasi Likelihood under the Independence model Criterion (QIC) X ² : 1.83	-	-	-	-	-

(Continues)

TABLE 2 (Continued)

Mortality prediction model	Validated? ^a		AUROC (95% CI) Development cohort ^b	Calibration Development cohort ^b	Overall performance Development cohort ^b	Type of validation cohort in original publication	AUROC (95% CI) Validation cohort	Calibration Validation cohort	Overall performance Validation cohort
	Internally	Externally							
COPE Duke et al ³⁶	-	+ Original publication	0.83 (0.83-0.84)	HL X ² : 23.1 (P < .01)	-	External validation dataset	0.83 (0.83-0.84)	HL X ² : 26.9 (P < .01)	-
PREDICT Ho et al ³⁷	+ Bootstrapping	+	-	-	-	Internal validation dataset	0.76 (0.75-0.77)	Calibration plot present	Nagelkerke's R ² : 0.255
High-Risk Selection System Iapichino et al ⁴⁶	+ Data splitting	-	0.81 (n.s.)	HL X ² : n.s. (P = .21)	-	Internal validation dataset	0.81 (n.s.)	HL X ² : n.s. (P = .22)	-
GV-SAPS II Liu et al ⁴⁷	-	+ Original publication	0.83 (0.81-0.84)	-	-	External validation dataset	0.82 (0.81-0.83)	-	-
MODS/NEMS Kao et al ³⁸	+ Bootstrapping	-	0.79 (n.s.)	-	-	Internal validation dataset	0.76 (n.s.)	HL X ² : 5.48 (P = .32) ^c	-
SMS-ICU Granholm et al ⁴⁸	+ Bootstrapping	+	0.72 (0.71-0.74)	HL X ² : 9.0 (P = .34) ^c	Nagelkerke's R ² : 0.191	Internal validation dataset	0.73 (n.s.)	Calibration slope: 0.99	Nagelkerke's R ² : 0.193
P-model Umegaki et al ³⁹	+ Cross-validation	-	0.87 (0.85-0.90)	HL X ² : 14.5 (P = .07)	-	Internal validation dataset	0.90 (0.88-0.92)	HL X ² : 13.5 (P = .10)	-
BCV model Huang et al ⁴⁰	+ Data splitting	-	0.79 (0.76-0.81)	HL X ² : 8.7 (P = .37)	-	Internal validation dataset	0.76 (0.71-0.81)	HL X ² : 11.1 (P = .19)	-
BCV/APACHE II model Huang et al ⁴⁰	+ Data splitting	-	0.80 (0.78-0.83)	HL X ² : 6.2 (P = .63)	-	Internal validation dataset	0.78 (0.73-0.83)	HL X ² : 5.4 (P = .72)	-
CREEK Stachon et al ⁴¹	+ Cross-validation	-	0.86 (n.s.)	HL C-statistic: 10.7 (P = .22) HL H-statistic: 10.1 (P = .26)	Brier score: 0.096	Internal validation dataset	0.832 (n.s.)	-	-
SAPS-R Viviani et al ⁴²	-	+ Original publication	-	-	-	External validation dataset	0.76 (n.s.)	-	-

(Continues)

TABLE 2 (Continued)

Mortality prediction model	Validated? ^a		AUROC (95% CI) Development cohort ^b	Calibration Development cohort ^b	Overall performance Development cohort ^b	Type of validation cohort in original publication	AUROC (95% CI) Validation cohort	Calibration Validation cohort	Overall performance Validation cohort
	Internally	Externally							
SAPS-E Viviani et al ⁴²	-	+ Original publication	-	-	-	External validation dataset	0.79 (n.s.)	-	-
25OHD Deyo-Charlson Comorbidity Index Mahato et al ⁴⁹	-	-	0.75 (0.67-0.83)	-	-	-	-	-	-
DELAWARE Stachon et al ⁴³	-	+ Original publication	0.86 (0.80-0.91)	HL statistic: n.s. (P = .28) Calibration plot present	-	External validation dataset	0.81 (0.75-0.87)	HL statistic: 0.44 (P = n.s.) Calibration plot present	-
Simplified Mortality Score Goag et al ⁴⁴	+	-	-	-	-	Internal validation dataset	0.58 (n.s.)	-	-

Abbreviations: ANZROD, Australian and New Zealand Risk Of Death; APACHE, Acute Physiology and Chronic Health Evaluation; APM, adductor pollicis muscle; AUROC, area under the receiving operating curves; BCV, Blood Cell Variability; CI, confidence interval; COPE, Critical Care Outcome Prediction Equation; CREEK, Critical Risk Evaluation by Early Keys; DELAWARE, Dense Laboratory Whole Blood Applied Risk Estimation; GV, glucose variability; HL, Hosmer-Lemeshow; ICNARC, Intensive Care National Audit Research Centre; ICU, intensive care unit; MMI, Multi-morbidity Index; MODS, Multiple Organs Dysfunctional Score; MPM, mortality prediction model; NEMS, Nine Equivalents Nursing Manpower use Score; NQF-ICOMMort, National Quality Forum ICU outcomes model (mortality); n.s., not specified; OASIS, Oxford Acute Severity of Illness Score; PREDICT, Predicted Risk, Existing Diseases and Intensive Care Therapy; RDW, red cell distribution width; SAPS, Simplified Acute Physiology Score; SMS-ICU, Simplified Mortality Score for the Intensive Care Unit; SUPPORT, Study to Understand Prognoses and Preferences for Outcomes and Risks of Treatments; TRIOS, Three-day Recalibrating ICU Outcomes.

^aCitations of original publications were screened on internal and/or external validation articles and shown as being present (+) or not present (-). When internal validation was present, the method of internal validation used in the original publication was presented. When external validation in the original publication was present, original publication was added in the column.

^bDevelopment cohort indicates the cohort in whom the prediction model was developed, sometimes also referred to as training cohort.

^cNot clear whether the value was derived from the development or validation dataset in the original publication, or value was derived from the development and validation dataset together.

^dNot clear whether this value is calculated for the 6-month mortality outcome or 12-month mortality.

^eNot clear whether the value was derived from the internal or external validation dataset in the original publication.

models (60%).^{14-17,21,22,24-26,28,30,32-36,38-41,43,46,48} Calibration plot was expressed for 12 prediction models (28%),^{13,15,20,24,28,30,31,33,35,37,43,48} and two prediction models (4.7%) presented the calibration slope value.^{30,48} Finally, one prediction model (2.3%) used the likelihood ratio test chi-squared value,²³ and one prediction model (2.3%) used the Quasi likelihood under the Independence Criterion.⁴⁵ In 11 prediction models (26%), calibration was not assessed.^{6,18,19,27,29,42,44,47,49}

Overall performance was expressed as the Brier score in eight mortality prediction models (19%),^{6,13,28-31,34,41} and as Nagelkerke's R^2 in two prediction models (4.7%).^{37,48}

4 | DISCUSSION

4.1 | Main findings

In this scoping review, we presented a contemporary overview of 43 mortality prediction models used in adult ICU patients in high-income countries. We found varying methodology, and the validation and performance of individual prediction models differ. Only 23 mortality prediction models of the 43 (53%) were externally validated. This overview provides a basis for head-to-head comparison of existing mortality prediction models through systematic external validation, with the ultimate goal to identify the most suitable prediction model for a certain cohort of patients.

4.2 | Summary of evidence

In previous literature, the maximum number of ICU mortality prediction models reviewed was 12,⁷ which is considerably less than the 43 prediction models identified by this review. Where we included all developed prediction models specifically designed to assess mortality, other reviews regarding ICU mortality prediction models focused mainly on commonly used models like the APACHE, SAPS and MPM,³⁻⁵ or identified models with different outcome than mortality (eg organ dysfunction) or disease- or organ-specific prognostic models.^{4,5,7,8} Additionally, only Siontis *et al* and Strand *et al* applied a systematic search to identify the models and discussed the validation of the models.^{5,8} Where we included all developed mortality prediction models, Strand *et al* did only include prediction models when the search for the specific scoring system yielded more than 50 citations.⁵ Siontis *et al.* conducted an evaluation of validated tools for hospitalized patients to predict all-cause mortality. However, their analysis included specific patient groups (eg heart or liver patients) rather than general ICU patients as included in the current review.⁸

Model performance is affected by the choice of outcome.^{31,50} Most mortality prediction models used hospital mortality as outcome measure.^{6,13-19,21,22,24,27,28,30-33,35,36,38,41-43,45,46} In general, longer fixed-time outcome measures used in some models^{20,24,25,29,31,37,39,40,44,45,47-49} are currently recommended.⁵¹ To elaborate, hospital mortality is dependent on discharge practices

and availability of post-ICU care, and is therefore a subjective measure. Furthermore, critical illness affects patients after hospital discharge.

The time span during which the mortality prediction models gathered their data varied from short (eg upon ICU admission or during the first initial hour of admission to the ICU) to long (eg during the first 24 hours of admission). Concerning complexity (time consumption) and missing data problems, it may be better in some situations to use a simpler model with less missing data than a more complex model built from a dataset with more missing data which achieves a slightly better performance.⁵² Longer collection periods may lead to more complete data, as incompleteness is often substantial for biochemical variables for patients with short-duration admissions (ie less than 24 hours). However, sampling rate affects predictions.⁵³ This limitation is considered less important in models with shorter data collection. Similarly, the treatments administered during the first 24 hours in the ICU obviously also affect predictions.

4.3 | Comparison of performance

We reported the performance of mortality prediction models in terms of discrimination, calibration and overall performance values. Direct comparison of prediction models predictive performances is not possible, as the development cohorts differed substantially from one another. As a consequence, prediction models cannot be considered interchangeable. Comparisons that are not done head-to-head in external samples independent of all models developed are at high risk of being misleading and may lead to inappropriate conclusions and resource use.¹²

Of 43, 26 (60%) mortality prediction models used the HL goodness-of-fit test for calibration.^{14-17,21,22,24-26,28,30,32-36,38-41,43,46,48} The HL test is commonly used, despite being frequently non-significant for small data cohorts and nearly always significant for large data cohorts.⁵⁴⁻⁵⁷ When only the HL test is reported without any calibration plot or table comparing predicted and observed outcome frequencies, inadequate information regarding calibration is provided.¹

Many ICU mortality prediction models are available and comparatively assessing their performance is a crucial task.⁴ In all, 25 articles compared the performance of the new model with existing models but used the same cohort of patients that was used in the development of the 'novel' model.^{6,13,14,16-18,20,22,24,26-30,32,34,40-47,49} This methodology is inherently biased in favor of the 'novel' model.^{54,57} Comparisons between prediction models should therefore only be executed in independent external validation samples not used to develop any of the models.

4.4 | Machine-learning algorithms

Mortality prediction models developed as an electronic model or derived from a machine-learning algorithm such as *AutoTriage*⁵⁸

were excluded in our manuscript since code and data availability are not requirements in all journals and this is necessary to reproduce the specific prediction model. However, code availability appears to be a rising trend.⁵⁹ Machine-learning-based prediction models seem to achieve increasingly higher accuracies and are becoming more dynamic,⁶⁰ although they still have to include a sufficiently large development and validation cohort to adequately assess performance and the risk of overfitting. However, a recent systematic review concluded that machine learning did not have superior performance over logistic regression for clinical prediction models.⁶¹

The association between mortality and variables may have changed since the original mortality prediction models were developed, for example, as a result of advancements in diagnostics and therapeutics.⁶² Mortality alone however is rarely the only outcome measure for interventional studies in ICU patients, and many trials, especially in sepsis, include an organ dysfunction score as part of ongoing patient assessment so that effects on morbidity can also be evaluated.³

Misuse of mortality prediction models can lead to inappropriate use of resources and potentially even mismanagement of patient care due to incorrect stratification.⁵⁷ Awareness of the differences in model design, the variance of predictions across different ICU settings and the effect of heterogeneity in populations are of utmost importance.

4.5 | Limitations

Some limitations of this study need to be addressed. First, having restricted our search to the period from 2008, relevant mortality prediction models might have been overlooked. Even though some of the most widely used mortality prediction models precede the screening period, we identified 16 prediction models that were published before 2008, but optimally searches have no time limit.⁶³ Second, we only included mortality prediction models originally developed for use in the ICU. Mortality prediction models not originally developed for mortality prediction in the ICU could still be valuable clinically. Third, in some original publications, it was unclear whether the presented discrimination, calibration and/or overall performance values were derived from the development cohort or from the validation dataset. We aimed to clarify these, but certain values might reflect another dataset from the original publication. Fourth, we only provided a systematic overview of all developed mortality prediction models in adult critically ill patients. We did not perform a systematic review of every retrieved model complete with all consecutive internal and external validations, as results from different external validations in different cohorts are not directly comparable due to differences in populations, case-mix and settings. We restricted the scope of this review to only identify whether internal or external validation had been performed as a measure of thoroughness of development of the identified models. For this reason, only screening of citations of the original articles was done to identify internal and/or external validation articles. Therefore, we should address that

our assessment on mortality prediction models not being internally and/or externally validated might be incomplete if validation in different publications was missed. A systematic search specifically designed for retrieving validation papers is advised when systematically reviewing the internal and external validations of mortality prediction models.⁶⁴

4.6 | Unanswered research questions

Although we retrieved many developed mortality prediction models that can be used as a step towards future head-to-head comparison, with the results of this scoping review it is not possible to make a recommendation on what mortality prediction models to use and it was not our intention to do so. External validation involving direct head-to-head comparisons in independent cohorts is needed to unravel the comparable performance of individual models. Although we provide a systematic overview of mortality prediction models and describe whether these were internally and/or externally validated, it was not desirable to give an overview of all external validations of the prediction models since this would require a specific search strategy for each model. Moreover, we would have liked to assess risk of bias using the recently developed PROBAST score.¹ However, this was not feasible because of the number of prediction models.

5 | FUTURE PERSPECTIVES

To identify the most suitable mortality prediction model for a certain patient cohort, ideally a head-to-head comparison of available models should be performed through systematic external validation using prospectively obtained datasets and appropriate statistical methods. The eventual aim will be to use this review to identify, update and implement the best performing mortality prediction models in daily practice. We are in the process of validating the found prediction models in independent contemporary cohorts to provide external validation of these models. Second, the process should be performed in different cohorts as heterogeneity of ICU patients exists on multiple levels, that is, patient level, hospital level, region and country level.⁶⁵ The best mortality prediction model in one setting is not necessarily the best performing prediction model in another setting. Third, it is worth mentioning that ICU patients have reduced long-term survival and impaired quality of life after ICU discharge compared to the general population.⁶⁶ Future research should also look at determinants of poor outcomes in ICU survivors to help guide long-term follow-up.⁶⁷

6 | CONCLUSIONS










In this review, 43 mortality prediction models have been studied. The validation and performance of individual prediction models

differ and the best prediction models for guiding clinical care and research is still to be established.

COMPETING INTERESTS/DISCLOSURES

AG and MHM were involved in the development of one of the mortality prediction models included. RGP reports shares in Evidencio BV, an online platform aiming to facilitate the creation, validation and implementation of clinical prediction models. Evidencio was not involved in the development of any of the prediction models mentioned nor is expecting to be affected financially by publication of this scoping review.

ORCID

Britt E. Keuning  <https://orcid.org/0000-0002-6222-7783>
 Thomas Kaufmann  <https://orcid.org/0000-0003-0589-8879>
 Renske Wiersema  <https://orcid.org/0000-0003-2413-2852>
 Anders Granholm  <https://orcid.org/0000-0001-5799-7655>
 Ville Pettilä  <https://orcid.org/0000-0002-3921-4423>
 Morten Hylander Møller  <https://orcid.org/0000-0002-6378-9673>
 Christian Fynbo Christiansen  <https://orcid.org/0000-0002-0727-953X>
 José Castela Forte  <https://orcid.org/0000-0001-9273-0702>
 Harold Snieder  <https://orcid.org/0000-0003-1949-2298>
 Frederik Keus  <https://orcid.org/0000-0003-1516-1475>
 Rick G. Pleijhuis  <https://orcid.org/0000-0003-4856-019X>
 Iwan C. C. van der Horst  <https://orcid.org/0000-0003-3891-8522>

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SUPPORTING INFORMATION

Additional supporting information may be found online in the Supporting Information section.

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